

REMARKS

A Notice of Allowance has been mailed from the USPTO for the above-referenced application. The Examiner contacted the undersigned by telephone indicating that Tables 5 and 6 were not sufficiently clear for printing the forthcoming Letters Patent. Applicants submit herewith Replacement Sheets containing Tables 5 and 6. No issue of new matter arises from this submission since the Replacement Sheets are merely clearer copies of the Tables 5 and 6 as filed.

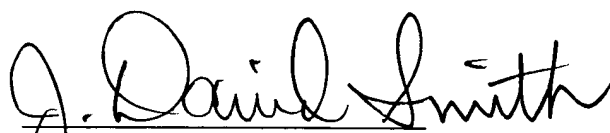
Fees

No fees are believed to be necessary in connection with this Amendment. However, if this is in error, authorization is hereby given to charge Deposit Account No. 11-1153 for any underpayment, or credit any overages.

Conclusion

If a discussion might be of assistance in resolving any issues, the Examiner is invited to telephone the undersigned.

Respectfully submitted,



J. David Smith
Attorney for Applicant(s)
Registration No. 39,839

KLAUBER & JACKSON
411 Hackensack Avenue
Hackensack, NJ 07601
(201) 487-5800

Table 5

Atomic Structure Coordinates of the
Free Form of the P/CAF Bromodomain

REMARK FILENAME: /biochem/chr18/BROMO_P/CAF/structures/1f8d/187.pdb									
REMARK Initial random number seed: 1342876511									
REMARK =====									
REMARK Bond, angles, improper, noe cdh									
REMARK =====									
REMARK num-deriv: 2.164156E-03, 0.365411, 50.3111, 41.9893E-02, 0.263503									
REMARK =====									
REMARK noe, cdh									
REMARK =====									
REMARK DATE: 22-Nov-98 06:51:33 created by user: jmm									
REMARK =====									
ATOM	1	CA	GLY	1	27.208	16.825	-6.349	1.00	0.00
ATOM	2	HA1	GLY	1	26.763	16.827	-5.365	1.00	0.00
ATOM	3	HA2	GLY	1	28.041	17.514	-6.357	1.00	0.00
ATOM	4	GLY	1	27.723	15.432	-6.650	1.00	0.00	0.00
ATOM	5	GLY	1	26.558	15.263	-7.435	1.00	0.00	0.00
ATOM	6	GLY	1	26.558	15.263	-7.435	1.00	0.00	0.00
ATOM	7	HA1	GLY	1	26.558	15.263	-7.435	1.00	0.00
ATOM	8	HA2	GLY	1	26.558	15.263	-7.435	1.00	0.00
ATOM	9	HA3	GLY	1	25.350	17.641	-6.843	1.00	0.00
ATOM	10	HA4	GLY	1	25.913	16.498	-7.955	1.00	0.00
ATOM	11	HA5	GLY	1	27.913	14.431	-6.024	1.00	0.00
ATOM	12	HA6	GLY	1	26.376	14.829	-5.409	1.00	0.00
ATOM	13	HA7	GLY	1	26.486	13.047	-6.726	1.00	0.00
ATOM	14	HA8	GLY	1	27.638	13.329	-4.193	1.00	0.00
ATOM	15	HA9	GLY	1	26.904	12.729	-4.193	1.00	0.00
ATOM	16	HA10	GLY	1	27.462	11.277	-5.017	1.00	0.00
ATOM	17	HA11	GLY	1	28.957	12.508	-4.319	1.00	0.00
ATOM	18	HA12	GLY	1	29.571	12.014	-4.982	1.00	0.00
ATOM	19	HA13	GLY	1	26.519	12.388	-4.834	1.00	0.00
ATOM	20	HA14	GLY	1	26.519	12.388	-4.834	1.00	0.00
ATOM	21	HA15	GLY	1	27.022	11.698	-8.183	1.00	0.00
ATOM	22	HA16	GLY	1	27.986	11.747	-8.353	1.00	0.00
ATOM	23	HA17	GLY	1	26.137	10.955	-9.106	1.00	0.00
ATOM	24	HA18	GLY	1	26.594	11.222	-10.553	1.00	0.00
ATOM	25	HA19	GLY	1	26.594	11.222	-10.553	1.00	0.00
ATOM	26	HA20	GLY	1	26.594	11.222	-10.553	1.00	0.00
ATOM	27	HA21	GLY	1	26.594	11.222	-10.553	1.00	0.00
ATOM	28	HA22	GLY	1	25.611	10.612	-11.561	1.00	0.00
ATOM	29	HA23	GLY	1	25.611	10.612	-11.561	1.00	0.00
ATOM	30	HA24	GLY	1	25.985	10.494	-12.900	1.00	0.00
ATOM	31	HA25	GLY	1	26.820	10.790	-13.130	1.00	0.00
ATOM	32	HA26	GLY	1	26.433	10.080	-11.420	1.00	0.00
ATOM	33	HA27	GLY	1	23.870	10.003	-10.501	1.00	0.00
ATOM	34	HA28	GLY	1	24.981	9.918	-11.537	1.00	0.00
ATOM	35	HA29	GLY	1	26.105	9.811	-7.536	1.00	0.00
ATOM	36	HA30	GLY	1	26.304	9.811	-6.890	1.00	0.00
ATOM	37	HA31	GLY	1	26.304	9.811	-6.890	1.00	0.00
ATOM	38	HA32	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	39	HA33	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	40	HA34	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	41	HA35	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	42	HA36	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	43	HA37	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	44	HA38	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	45	HA39	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	46	HA40	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	47	HA41	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	48	HA42	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	49	HA43	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	50	HA44	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	51	HA45	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	52	HA46	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	53	HA47	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	54	HA48	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	55	HA49	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	56	HA50	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	57	HA51	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	58	HA52	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	59	HA53	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	60	HA54	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	61	HA55	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	62	HA56	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	63	HA57	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	64	HA58	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	65	HA59	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	66	HA60	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	67	HA61	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	68	HA62	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	69	HA63	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	70	HA64	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	71	HA65	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	72	HA66	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	73	HA67	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	74	HA68	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	75	HA69	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	76	HA70	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	77	HA71	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	78	HA72	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	79	HA73	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	80	HA74	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	81	HA75	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	82	HA76	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	83	HA77	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	84	HA78	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	85	HA79	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	86	HA80	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	87	HA81	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	88	HA82	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	89	HA83	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	90	HA84	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	91	HA85	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	92	HA86	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	93	HA87	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	94	HA88	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	95	HA89	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	96	HA90	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	97	HA91	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	98	HA92	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	99	HA93	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	100	HA94	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	101	HA95	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	102	HA96	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	103	HA97	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	104	HA98	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	105	HA99	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	106	HA100	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	107	HA101	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	108	HA102	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	109	HA103	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	110	HA104	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	111	HA105	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	112	HA106	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	113	HA107	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	114	HA108	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	115	HA109	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	116	HA110	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	117	HA111	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	118	HA112	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	119	HA113	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	120	HA114	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	121	HA115	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	122	HA116	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	123	HA117	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	124	HA118	GLY	1	26.213	9.459	-8.817	1.00	0.00
ATOM	125	HA119	GLY						

REPLACEMENT SHEET

ATOM	255	HQ1 THH	17	10.098	1.223	6.144	1.00	0.00	ATOM	349	HQ23 LBU	22	0.803	-4.924	3.008	1.00	0.00	ATOM	443	ND1 H1S	28	-4.254	1.759	13.538	1.00	0.00
ATOM	256	CO2 THH	17	7.971	0.832	4.783	1.00	0.00	ATOM	350	C LBU	22	1.493	-5.568	8.214	1.00	0.00	ATOM	444	ND1 H1S	28	-4.081	1.459	11.558	1.00	0.00
ATOM	257	HQ21 THH	17	7.078	0.224	4.720	1.00	0.00	ATOM	351	O LBU	22	2.546	-5.862	8.525	1.00	0.00	ATOM	445	ND2 H1S	28	-4.569	1.777	10.377	1.00	0.00
ATOM	258	HQ22 THH	17	8.088	1.395	3.867	1.00	0.00	ATOM	352	N LBU	22	0.310	-5.868	8.971	1.00	0.00	ATOM	446	ND3 H1S	28	-4.665	1.477	9.340	1.00	0.00
ATOM	259	HQ23 THH	17	7.883	1.513	5.616	1.00	0.00	ATOM	353	HN GLN	23	3.447	-5.617	8.666	1.00	0.00	ATOM	447	ND4 H1S	28	-4.537	3.006	12.191	1.00	0.00
ATOM	260	THH	17	7.883	1.513	5.616	1.00	0.00	ATOM	354	CA GLN	23	2.402	-5.598	10.266	1.00	0.00	ATOM	448	ND5 H1S	28	-4.587	3.006	12.191	1.00	0.00
ATOM	261	O THH	17	6.752	-1.785	8.821	1.00	0.00	ATOM	355	HN GLN	23	2.006	-5.756	9.989	1.00	0.00	ATOM	449	ND6 H1S	28	-4.719	3.042	10.886	1.00	0.00
ATOM	262	N LBU	18	7.574	-2.569	4.697	1.00	0.00	ATOM	356	CA GLN	23	3.568	-6.635	12.006	1.00	0.00	ATOM	450	ND7 H1S	28	-4.535	3.046	10.886	1.00	0.00
ATOM	263	HN LBU	18	8.319	-3.578	4.083	1.00	0.00	ATOM	357	HN GLN	23	3.568	-6.635	12.006	1.00	0.00	ATOM	451	ND8 H1S	28	-4.535	3.046	10.886	1.00	0.00
ATOM	264	CA LBU	18	6.369	-3.302	4.310	1.00	0.00	ATOM	358	HN GLN	23	4.528	-5.842	10.640	1.00	0.00	ATOM	452	ND9 H1S	28	-6.982	-1.724	11.800	1.00	0.00
ATOM	265	HN LBU	18	5.555	-2.597	4.289	1.00	0.00	ATOM	359	HN GLN	23	4.528	-5.842	10.640	1.00	0.00	ATOM	453	ND10 H1S	28	-6.982	-1.724	11.800	1.00	0.00
ATOM	266	CA LBU	18	6.369	-3.302	4.310	1.00	0.00	ATOM	360	HN GLN	23	4.528	-5.842	10.640	1.00	0.00	ATOM	454	ND11 H1S	28	-6.982	-1.724	11.800	1.00	0.00
ATOM	267	HN LBU	18	5.555	-2.597	4.289	1.00	0.00	ATOM	361	HN GLN	23	4.528	-5.842	10.640	1.00	0.00	ATOM	455	ND12 H1S	28	-6.982	-1.724	11.800	1.00	0.00
ATOM	268	CA LBU	18	6.369	-3.302	4.310	1.00	0.00	ATOM	362	HN GLN	23	4.528	-5.842	10.640	1.00	0.00	ATOM	456	ND13 H1S	28	-6.982	-1.724	11.800	1.00	0.00
ATOM	269	HN LBU	18	5.555	-2.597	4.289	1.00	0.00	ATOM	363	HN GLN	23	4.528	-5.842	10.640	1.00	0.00	ATOM	457	ND14 H1S	28	-6.982	-1.724	11.800	1.00	0.00
ATOM	270	HN LBU	18	6.964	-2.940	1.871	1.00	0.00	ATOM	364	HN GLN	23	3.958	-10.308	10.517	1.00	0.00	ATOM	458	ND15 H1S	28	-8.012	2.021	14.552	1.00	0.00
ATOM	271	CD1 LBU	18	6.954	-3.607	0.510	1.00	0.00	ATOM	365	HN GLN	23	3.958	-10.308	10.517	1.00	0.00	ATOM	459	ND16 H1S	28	-8.012	2.021	14.552	1.00	0.00
ATOM	272	CD1 LBU	18	7.945	-3.607	0.510	1.00	0.00	ATOM	366	HN GLN	23	4.439	-10.275	9.658	1.00	0.00	ATOM	460	ND17 H1S	28	-7.670	0.341	15.809	1.00	0.00
ATOM	273	HN LBU	18	6.480	-2.894	-0.236	1.00	0.00	ATOM	367	C GLN	23	1.374	-5.904	11.151	1.00	0.00	ATOM	461	ND18 H1S	28	-7.180	-0.560	15.463	1.00	0.00
ATOM	274	HN LBU	18	6.480	-2.894	-0.236	1.00	0.00	ATOM	368	C GLN	23	0.642	-5.604	11.646	1.00	0.00	ATOM	462	ND19 H1S	28	-6.929	1.021	16.203	1.00	0.00
ATOM	275	HN LBU	18	6.270	-4.442	1.524	1.00	0.00	ATOM	369	N GLN	24	1.562	-4.843	11.044	1.00	0.00	ATOM	463	ND20 H1S	28	-8.364	-0.244	16.997	1.00	0.00
ATOM	276	HN LBU	18	5.118	-1.943	1.430	1.00	0.00	ATOM	370	N GLN	24	2.053	-4.937	11.317	1.00	0.00	ATOM	464	ND21 H1S	28	-8.364	-0.244	16.997	1.00	0.00
ATOM	277	HN LBU	18	5.118	-1.943	1.430	1.00	0.00	ATOM	371	N GLN	24	2.053	-4.937	11.317	1.00	0.00	ATOM	465	ND22 H1S	28	-8.364	-0.244	16.997	1.00	0.00
ATOM	278	HN LBU	18	5.118	-1.943	1.430	1.00	0.00	ATOM	372	N GLN	24	0.434	-4.586	13.142	1.00	0.00	ATOM	466	ND23 H1S	28	-8.870	-0.866	15.613	1.00	0.00
ATOM	279	HN LBU	18	6.037	-4.376	5.349	1.00	0.00	ATOM	373	HN GLN	24	1.369	-2.679	13.260	1.00	0.00	ATOM	467	ND24 H1S	28	-10.363	-0.935	17.267	1.00	0.00
ATOM	280	C LBU	18	4.961	-4.376	5.349	1.00	0.00	ATOM	374	HN GLN	24	2.330	-2.679	13.260	1.00	0.00	ATOM	468	ND25 H1S	28	-8.607	1.069	12.032	1.00	0.00
ATOM	281	N LBU	19	6.956	-5.310	5.942	1.00	0.00	ATOM	375	HN GLN	24	1.974	-2.754	13.659	1.00	0.00	ATOM	469	ND26 H1S	28	-9.794	1.378	12.032	1.00	0.00
ATOM	282	N LBU	19	6.753	-5.310	5.942	1.00	0.00	ATOM	376	HN GLN	24	1.974	-2.754	13.659	1.00	0.00	ATOM	470	ND27 H1S	28	-7.711	1.378	12.032	1.00	0.00
ATOM	283	CA LBU	19	5.946	-6.295	6.158	1.00	0.00	ATOM	377	HN GLN	24	0.960	-3.077	11.689	1.00	0.00	ATOM	471	ND28 H1S	28	-6.788	1.056	11.196	1.00	0.00
ATOM	284	HA LBU	19	8.029	-7.225	6.535	1.00	0.00	ATOM	378	CD GLN	24	2.702	-0.638	12.200	1.00	0.00	ATOM	472	ND29 H1S	28	-9.148	2.199	9.989	1.00	0.00
ATOM	285	CB LBU	19	8.522	-7.243	5.474	1.00	0.00	ATOM	379	CD GLN	24	3.210	-0.671	13.320	1.00	0.00	ATOM	473	ND30 H1S	28	-7.375	3.385	9.974	1.00	0.00
ATOM	286	HN LBU	19	7.748	-8.232	6.897	1.00	0.00	ATOM	380	HN GLN	24	3.082	0.227	11.265	1.00	0.00	ATOM	474	ND31 H1S	28	-6.453	3.289	9.324	1.00	0.00
ATOM	287	HN LBU	19	9.033	-6.730	6.897	1.00	0.00	ATOM	381	HN GLN	24	3.795	0.862	11.468	1.00	0.00	ATOM	475	ND32 H1S	28	-8.013	4.090	9.324	1.00	0.00
ATOM	288	HN LBU	19	9.033	-6.730	6.897	1.00	0.00	ATOM	382	HN GLN	24	-0.929	-3.421	11.254	1.00	0.00	ATOM	476	ND33 H1S	28	-7.071	3.890	11.196	1.00	0.00
ATOM	289	HN LBU	19	9.033	-6.730	6.897	1.00	0.00	ATOM	383	HN GLN	24	-0.929	-3.421	11.254	1.00	0.00	ATOM	477	ND34 H1S	28	-7.071	3.890	11.196	1.00	0.00
ATOM	290	HN LBU	19	9.033	-6.730	6.897	1.00	0.00	ATOM	384	HN GLN	24	-0.929	-3.421	11.254	1.00	0.00	ATOM	478	ND35 H1S	28	-7.071	3.890	11.196	1.00	0.00
ATOM	291	CD LBU	19	10.412	-7.319	7.429	1.00	0.00	ATOM	385	C GLN	24	-0.577	-2.936	10.456	1.00	0.00	ATOM	479	ND36 H1S	28	-8.056	1.145	7.611	1.00	0.00
ATOM	292	HN LBU	19	10.519	-7.440	6.365	1.00	0.00	ATOM	386	N VAL	25	-0.577	-2.936	10.456	1.00	0.00	ATOM	480	ND37 H1S	28	-8.056	1.145	7.611	1.00	0.00
ATOM	293	HN LBU	19	11.156	-6.643	7.823	1.00	0.00	ATOM	387	N VAL	25	-0.286	-2.811	10.407	1.00	0.00	ATOM	481	ND38 H1S	28	-7.089	0.039	8.962	1.00	0.00
ATOM	294	CE LBU	19	10.567	-8.671	8.106	1.00	0.00	ATOM	388	N VAL	25	-1.769	-2.477	9.743	1.00	0.00	ATOM	482	ND39 H1S	28	-6.726	-0.224	9.860	1.00	0.00
ATOM	295	HN LBU	19	9.987	-8.671	8.106	1.00	0.00	ATOM	389	N VAL	25	-2.139	-2.597	9.743	1.00	0.00	ATOM	483	ND40 H1S	28	-6.726	-0.224	9.860	1.00	0.00
ATOM	296	HN LBU	19	9.987	-8.671	8.106	1.00	0.00	ATOM	390	N VAL	25	-2.139	-2.597	9.743	1.00	0.00	ATOM	484	ND41 H1S	28	-6.726	-0.224	9.860	1.00	0.00
ATOM	297	HN LBU	19	11.988	-8.538	9.117	1.00	0.00	ATOM	391	N VAL	25	-2.139	-2.597	9.743	1.00	0.00	ATOM	485	ND42 H1S	28	-6.726	-0.224	9.860	1.00	0.00
ATOM	298	HN LBU	19	12.044	-10.147	8.010	1.00	0.00	ATOM	392	CD VAL	25	-2.139	-2.597	9.743	1.00	0.00	ATOM	486	ND43 H1S	28	-6.726	-0.224	9.860	1.00	0.00
ATOM	299	HN LBU	19	12.418	-8.655	9.057	1.00	0.00	ATOM	393	CD VAL	25	-0.774	-0.103	7.888	1.00	0.00	ATOM	487	ND44 H1S	28	-5.885	-2.407	9.054	1.00	0.00
ATOM	300	HN LBU	19	12.526	-8.655	7.380	1.00	0.00	ATOM	394	HN VAL	25	0.434	-1.358	7.591	1.00	0.00	ATOM	488	ND45 H1S	28	-4.765	-1.111	8.629	1.00	0.00
ATOM	301	C LBU	19	6.349	-5.772	7.863	1.00	0.00	ATOM	395	HN VAL	25	0.024	-0.930	7.424	1.00	0.00	ATOM	489	ND46 H1S	28	-7.906	-1.726	6.480	1.00	0.00
ATOM	302	C LBU	19	5.624	-6.390	8.446	1.00	0.00	ATOM	396	HN VAL	25	-1.964	-3.301	7.424	1.00	0.00	ATOM	490	ND47 H1S	28	-7.906	-1.726	6.480	1.00	0.00
ATOM	303	C LBU	19	5.624	-6.390	8.446	1.00	0.00	ATOM	397	HN VAL	25	-1.964	-3.301	7.424	1.00	0.00	ATOM	491	ND48 H1S	28	-7.906	-1.726	6.480	1.00	0.00
ATOM	304	HN LBU	20	7.346	-4.052	7.410	1.00	0.00	ATOM	398	HN VAL	25	-1.964	-3.301	7.424	1.00	0.00	ATOM	492	ND49 H1S	28	-7.906	-1.726	6.480	1.00	0.00
ATOM	305	HA LBU	20	6.482	-4.841	9.316	1.00	0.00	ATOM	399	HN VAL	25	-0.462	-2.982	6.511	1.00	0.00	ATOM	493	ND50 H1S	28	-9.946	7.971	1.00	0.00	
ATOM	306	HA LBU	20	6.482	-4.841	9.316	1.00	0.00	ATOM	400	C VAL															

REPLACEMENT SHEET

ATOM	537	CDL PHE	34	-6.937	0.754	3.561	1.00	0.00	ATOM	631	C LYS	39	-14.389	-9.553	-5.459	1.00	0.00	ATOM	725	CDL TYR	46	-10.532	-1.931	-7.399	1.00	0.00
ATOM	538	HDL PHE	34	-6.938	0.503	3.562	1.00	0.00	ATOM	632	C LYS	39	-13.886	-10.337	-7.297	1.00	0.00	ATOM	726	HDL TYR	46	-9.944	-2.650	-8.199	1.00	0.00
ATOM	539	CDZ PHE	34	-5.300	0.255	4.946	1.00	0.00	ATOM	633	NH ARG	40	-15.448	-8.206	-6.079	1.00	0.00	ATOM	727	CDZ TYR	46	-12.432	-0.609	-8.995	1.00	0.00
ATOM	540	HDL PHE	34	-6.111	-0.459	5.373	1.00	0.00	ATOM	634	NH ARG	40	-15.810	-8.206	-6.079	1.00	0.00	ATOM	728	HDL TYR	46	-13.194	0.029	-8.724	1.00	0.00
ATOM	541	CEL PHE	34	-7.069	2.077	3.868	1.00	0.00	ATOM	635	CA ARG	40	-16.078	-8.826	-8.074	1.00	0.00	ATOM	729	CEL TYR	46	-10.487	-1.522	-6.486	1.00	0.00
ATOM	542	CEL PHE	34	-7.761	2.789	3.443	1.00	0.00	ATOM	636	HA ARG	40	-15.347	-8.504	-8.800	1.00	0.00	ATOM	730	CEL TYR	46	-9.725	-1.952	-5.593	1.00	0.00
ATOM	543	CEL PHE	34	-7.368	1.536	5.459	1.00	0.00	ATOM	637	CA ARG	40	-16.534	-10.246	-8.417	1.00	0.00	ATOM	731	CEL TYR	46	-12.501	0.007	-6.783	1.00	0.00
ATOM	544	CEL PHE	34	-7.368	1.536	5.459	1.00	0.00	ATOM	638	HA ARG	40	-15.926	-10.197	-8.417	1.00	0.00	ATOM	732	CEL TYR	46	-12.501	0.007	-6.783	1.00	0.00
ATOM	545	CZ PHE	34	-6.252	2.450	4.917	1.00	0.00	ATOM	639	HA ARG	40	-17.946	-10.197	-8.417	1.00	0.00	ATOM	733	CEL TYR	46	-11.139	-0.143	-4.756	1.00	0.00
ATOM	546	HZ PHE	34	-6.304	3.453	5.314	1.00	0.00	ATOM	640	CO ARG	40	-17.189	-10.968	-7.216	1.00	0.00	ATOM	734	CO TYR	46	-10.892	-0.871	-4.339	1.00	0.00
ATOM	547	C PHE	34	-7.400	-3.624	3.053	1.00	0.00	ATOM	641	HDL ARG	40	-16.612	-10.809	-6.363	1.00	0.00	ATOM	735	HI TYR	46	-11.427	-0.671	-12.356	1.00	0.00
ATOM	548	O PHE	34	-7.293	-4.349	2.065	1.00	0.00	ATOM	642	HOZ ARG	40	-17.305	-12.462	-7.513	1.00	0.00	ATOM	736	O TYR	46	-11.940	-0.764	-11.971	1.00	0.00
ATOM	549	NH PHE	35	-7.543	-4.100	4.285	1.00	0.00	ATOM	643	CO ARG	40	-16.418	-12.788	-8.038	1.00	0.00	ATOM	737	O TYR	46	-10.120	-0.506	-12.072	1.00	0.00
ATOM	550	HN PHE	35	-7.653	-3.446	5.035	1.00	0.00	ATOM	644	HOZ ARG	40	-17.912	-12.160	-5.044	1.00	0.00	ATOM	738	NH TYR	47	-9.772	-0.440	-11.072	1.00	0.00
ATOM	551	HN PHE	35	-7.653	-3.446	5.035	1.00	0.00	ATOM	645	HN ARG	40	-17.912	-12.160	-5.044	1.00	0.00	ATOM	739	HN TYR	47	-9.772	-0.440	-11.072	1.00	0.00
ATOM	552	CB MET	35	-6.624	-5.948	4.281	1.00	0.00	ATOM	646	HN ARG	40	-17.912	-12.160	-5.044	1.00	0.00	ATOM	740	HN TYR	47	-9.772	-0.440	-11.072	1.00	0.00
ATOM	553	CB MET	35	-7.868	-5.802	6.021	1.00	0.00	ATOM	647	CB ARG	40	-17.683	-13.735	-5.465	1.00	0.00	ATOM	741	CB TYR	47	-8.188	-0.506	-12.860	1.00	0.00
ATOM	554	HBI MET	35	-8.472	-4.998	6.413	1.00	0.00	ATOM	648	CZ ARG	40	-17.235	-14.537	-6.197	1.00	0.00	ATOM	742	CB TYR	47	-9.433	-1.567	-14.821	1.00	0.00
ATOM	555	HBI MET	35	-8.415	-6.729	6.110	1.00	0.00	ATOM	649	HN ARG	40	-16.898	-15.225	-7.219	1.00	0.00	ATOM	743	HBI TYR	47	-10.524	-1.285	-15.460	1.00	0.00
ATOM	556	CO MET	35	-6.609	-5.910	6.868	1.00	0.00	ATOM	650	HN ARG	40	-16.794	-14.759	-8.157	1.00	0.00	ATOM	744	HBI TYR	47	-9.839	-1.748	-14.743	1.00	0.00
ATOM	557	CO MET	35	-6.609	-5.910	6.868	1.00	0.00	ATOM	651	HN ARG	40	-16.794	-14.759	-8.157	1.00	0.00	ATOM	745	CO TYR	47	-9.839	-1.748	-14.743	1.00	0.00
ATOM	558	HOZ MET	35	-5.714	-5.267	7.750	1.00	0.00	ATOM	652	HN ARG	40	-17.912	-12.160	-5.044	1.00	0.00	ATOM	746	HOZ TYR	47	-10.783	-3.715	-14.055	1.00	0.00
ATOM	559	SD MET	35	-6.296	-7.591	7.437	1.00	0.00	ATOM	653	HN ARG	40	-17.912	-12.160	-5.044	1.00	0.00	ATOM	747	HOZ TYR	47	-10.783	-3.715	-14.055	1.00	0.00
ATOM	560	CE MET	35	-8.484	-8.011	6.470	1.00	0.00	ATOM	654	HN ARG	40	-17.912	-12.160	-5.044	1.00	0.00	ATOM	748	CDZ TYR	47	-9.290	-3.229	-13.290	1.00	0.00
ATOM	561	CEL MET	35	-4.248	-8.726	7.013	1.00	0.00	ATOM	655	C ARG	40	-17.220	-16.147	-4.976	1.00	0.00	ATOM	749	CDZ TYR	47	-8.572	-2.574	-11.818	1.00	0.00
ATOM	562	HEZ MET	35	-4.268	-7.119	6.285	1.00	0.00	ATOM	656	O ARG	40	-17.220	-16.147	-4.976	1.00	0.00	ATOM	750	CEL TYR	47	-11.124	-4.899	-13.916	1.00	0.00
ATOM	563	HEZ MET	35	-5.157	-8.441	5.529	1.00	0.00	ATOM	657	O TYR	41	-18.081	-7.910	-7.065	1.00	0.00	ATOM	751	CEL TYR	47	-11.843	-5.550	-13.971	1.00	0.00
ATOM	564	O MET	35	-6.656	-6.134	6.574	1.00	0.00	ATOM	658	HN TYR	41	-17.886	-8.510	-6.332	1.00	0.00	ATOM	752	CEL TYR	47	-9.659	-4.411	-11.674	1.00	0.00
ATOM	565	NH MET	36	-8.219	-7.049	7.251	1.00	0.00	ATOM	659	HN TYR	41	-19.424	-8.751	-9.571	1.00	0.00	ATOM	753	HEZ TYR	47	-10.215	-4.619	-11.525	1.00	0.00
ATOM	566	NH MET	36	-7.255	-7.208	7.251	1.00	0.00	ATOM	660	HN TYR	41	-19.424	-8.751	-9.571	1.00	0.00	ATOM	754	HEZ TYR	47	-10.215	-4.619	-11.525	1.00	0.00
ATOM	567	HN MET	36	-9.128	-7.729	1.929	1.00	0.00	ATOM	661	CB TYR	41	-20.418	-7.806	-6.335	1.00	0.00	ATOM	755	HI TYR	47	-10.934	-6.431	-11.673	1.00	0.00
ATOM	568	CA MET	36	-8.598	-8.569	1.409	1.00	0.00	ATOM	662	CB TYR	41	-20.418	-7.806	-6.335	1.00	0.00	ATOM	756	HI TYR	47	-10.934	-6.431	-11.673	1.00	0.00
ATOM	569	HA MET	36	-10.371	-8.248	2.562	1.00	0.00	ATOM	663	COI TYR	41	-21.586	-7.003	-6.313	1.00	0.00	ATOM	757	C TYR	47	-9.318	0.917	-13.916	1.00	0.00
ATOM	570	CB MET	36	-11.151	-7.504	2.496	1.00	0.00	ATOM	664	HOI TYR	41	-21.499	-6.331	-5.634	1.00	0.00	ATOM	758	O TYR	47	-8.350	1.686	-13.992	1.00	0.00
ATOM	571	HBZ MET	36	-10.371	-7.504	2.496	1.00	0.00	ATOM	665	COZ TYR	41	-20.139	-8.255	-4.935	1.00	0.00	ATOM	759	NH MET	48	-10.520	1.184	-14.452	1.00	0.00
ATOM	572	CO MET	36	-10.907	-9.553	1.996	1.00	0.00	ATOM	666	HN TYR	41	-19.373	-8.020	-4.325	1.00	0.00	ATOM	760	HN MET	48	-11.227	0.529	-14.452	1.00	0.00
ATOM	573	CO MET	36	-10.907	-9.553	1.996	1.00	0.00	ATOM	667	HN TYR	41	-19.373	-8.020	-4.325	1.00	0.00	ATOM	761	HN MET	48	-11.227	0.529	-14.452	1.00	0.00
ATOM	574	HOZ MET	36	-11.473	-9.318	1.102	1.00	0.00	ATOM	668	HOZ TYR	41	-21.045	-8.644	-4.477	1.00	0.00	ATOM	762	HA TYR	48	-10.101	2.441	-16.037	1.00	0.00
ATOM	575	HOZ MET	36	-11.473	-9.318	1.102	1.00	0.00	ATOM	669	C TYR	41	-21.045	-8.644	-4.477	1.00	0.00	ATOM	763	CB MET	48	-12.309	2.478	-11.648	1.00	0.00
ATOM	576	CD MET	36	-11.806	-10.289	2.972	1.00	0.00	ATOM	670	O TYR	41	-19.562	-8.751	-6.318	1.00	0.00	ATOM	764	HBZ MET	48	-12.854	2.529	-11.784	1.00	0.00
ATOM	577	CD MET	36	-11.522	-11.469	3.566	1.00	0.00	ATOM	671	NH MET	42	-17.969	-8.915	-5.254	1.00	0.00	ATOM	765	HBZ MET	48	-12.343	3.368	-16.344	1.00	0.00
ATOM	578	CDZ MET	36	-12.794	-9.685	4.440	1.00	0.00	ATOM	672	HN MET	42	-17.911	-8.776	-5.156	1.00	0.00	ATOM	766	CB MET	48	-12.621	1.276	-16.482	1.00	0.00
ATOM	579	CDZ MET	36	-12.794	-9.685	4.440	1.00	0.00	ATOM	673	HN MET	42	-17.911	-8.776	-5.156	1.00	0.00	ATOM	767	CB MET	48	-12.621	1.276	-16.482	1.00	0.00
ATOM	580	O MET	36	-10.414	-5.939	6.970	1.00	0.00	ATOM	674	HN MET	42	-16.958	-8.524	-2.144	1.00	0.00	ATOM	768	HOI MET	48	-10.136	3.439	-14.992	1.00	0.00
ATOM	581	NH MET	37	-9.913	-6.939	-0.486	1.00	0.00	ATOM	675	CA MET	42	-15.513	-5.015	-3.864	1.00	0.00	ATOM	769	HOI MET	48	-10.136	3.439	-14.992	1.00	0.00
ATOM	582	CA MET	37	-9.200	-6.075	-1.444	1.00	0.00	ATOM	676	HBZ MET	42	-15.513	-5.015	-3.864	1.00	0.00	ATOM	770	CEL MET	48	-14.856	1.919	-15.994	1.00	0.00
ATOM	583	HA MET	37	-8.087	-5.425	-1.386	1.00	0.00	ATOM	677	HBZ MET	42	-16.561	-4.556	-3.279	1.00	0.00	ATOM	771	CEL MET	48	-10.454	0.433	-17.594	1.00	0.00
ATOM	584	CB MET	37	-7.650	-5.315	-3.021	1.00	0.00	ATOM	678	CB MET	42	-16.561	-4.556	-3.279	1.00	0.00	ATOM	772	O MET	48	-10.454	0.433	-17.594	1.00	0.00
ATOM	585	HBZ MET	37	-7.087	-7.224	-1.976	1.00	0.00	ATOM	679	HOZ MET	42	-17.313	-5.054	-1.482	1.00	0.00	ATOM	773	O MET	48	-10.136	3.439	-14.992	1.00	0.00
ATOM	586	HOZ MET	37	-6.627	-7.957	-2.522	1.00	0.00	ATOM	680	CA MET	42	-14.293	-3.913	-1.760	1.00	0.00	ATOM	774	CA MET	48	-10.261	4.545	-12.055	1.00	0.00
ATOM	587	CA MET	37	-6.338	-6.570	-1.455	1.00	0.00	ATOM	681	HOZ MET	42	-15.169	-5.300	-0.297	1.00	0.00	ATOM	775	CA MET	48	-9.905	5.392	-12.624	1.00	0.00
ATOM	588	HOZ MET	37	-7.857	-7.904	-0.784	1.00	0.00	ATOM	682	COZ MET	42	-17.072	-3.628	-5.249	1.00	0.00	ATOM	776	CA MET	48	-11.5				

REPLACEMENT SHEET

ATOM	819	CD	ARG	51	-2.313	4.849	-11.240	1.00	0.00	ATOM	913	CG	LVS	57	-6.542	-10.325	2.520	1.00	0.00	ATOM	1007	C	ARG	62	4.720	-11.542	-0.664	1.00	0.00
ATOM	820	HD	ARG	51	-2.996	5.010	-10.312	1.00	0.00	ATOM	914	HOL	LVS	57	-6.160	-10.141	1.629	1.00	0.00	ATOM	1008	C	ARG	62	5.781	-12.265	-0.576	1.00	0.00
ATOM	821	HD	ARG	51	-1.566	5.493	-11.246	1.00	0.00	ATOM	915	CD	LVS	57	-6.622	-10.954	2.517	1.00	0.00	ATOM	1009	N	LEU	63	4.700	-10.364	-0.052	1.00	0.00
ATOM	822	NE	ARG	51	-0.275	5.184	-12.385	1.00	0.00	ATOM	916	CD	LVS	57	-6.049	-11.720	3.735	1.00	0.00	ATOM	1010	N	LEU	63	3.882	-9.835	-0.149	1.00	0.00
ATOM	823	NE	ARG	51	-4.217	5.390	-12.212	1.00	0.00	ATOM	917	HOL	LVS	57	-5.944	-11.047	3.918	1.00	0.00	ATOM	1011	CA	LEU	63	5.792	-9.853	0.717	1.00	0.00
ATOM	824	CE	ARG	51	-2.832	5.226	-13.698	1.00	0.00	ATOM	918	HD	LVS	57	-5.090	-12.157	3.498	1.00	0.00	ATOM	1012	HA	LEU	63	6.363	-9.663	0.168	1.00	0.00
ATOM	825	CE	ARG	51	-1.332	4.955	-13.904	1.00	0.00	ATOM	919	CE	LVS	57	-7.017	-12.831	4.112	1.00	0.00	ATOM	1013	HB	LEU	63	5.388	-8.546	1.490	1.00	0.00
ATOM	826	HLL	ARG	51	-1.332	4.955	-13.904	1.00	0.00	ATOM	920	CE	LVS	57	-7.017	-12.831	4.112	1.00	0.00	ATOM	1014	HB	LEU	63	5.388	-8.546	1.490	1.00	0.00
ATOM	827	HLL	ARG	51	-1.332	4.955	-13.904	1.00	0.00	ATOM	921	NE	LVS	57	-6.621	-10.557	3.751	1.00	0.00	ATOM	1015	HB	LEU	63	4.460	-6.821	1.982	1.00	0.00
ATOM	828	NH2	ARG	51	-3.659	5.539	-14.666	1.00	0.00	ATOM	922	NE	LVS	57	-7.223	-12.915	5.585	1.00	0.00	ATOM	1016	CE	LEU	63	6.413	-7.984	2.436	1.00	0.00
ATOM	829	NH2	ARG	51	-3.323	5.569	-15.568	1.00	0.00	ATOM	923	H21	LVS	57	-6.363	-12.609	5.861	1.00	0.00	ATOM	1017	HG	LEU	63	7.441	-8.747	2.663	1.00	0.00
ATOM	830	CD	ARG	51	-4.617	5.746	-14.430	1.00	0.00	ATOM	924	H22	LVS	57	-7.441	-12.302	5.871	1.00	0.00	ATOM	1018	CD	LEU	63	7.148	-6.864	1.825	1.00	0.00
ATOM	831	C	ARG	51	-1.864	0.354	-11.056	1.00	0.00	ATOM	925	H23	LVS	57	-8.013	-12.302	5.871	1.00	0.00	ATOM	1019	HD1	LEU	63	6.457	-5.994	1.672	1.00	0.00
ATOM	832	C	ARG	51	-4.729	-0.139	-12.064	1.00	0.00	ATOM	926	C	LVS	57	-3.895	-10.144	1.617	1.00	0.00	ATOM	1020	H21	LEU	63	7.579	-7.097	0.877	1.00	0.00
ATOM	833	N	SER	52	-3.310	0.403	-9.751	1.00	0.00	ATOM	927	N	THR	58	-4.014	-9.716	0.370	1.00	0.00	ATOM	1021	HD2	LEU	63	5.741	-7.509	1.322	1.00	0.00
ATOM	834	HA	SER	52	-3.307	-1.652	-9.751	1.00	0.00	ATOM	928	N	THR	58	-4.014	-9.716	0.370	1.00	0.00	ATOM	1022	HD2	LEU	63	5.741	-7.509	1.322	1.00	0.00
ATOM	835	HA	SER	52	-3.766	-2.101	-10.603	1.00	0.00	ATOM	929	HA	THR	58	-4.551	-8.913	0.201	1.00	0.00	ATOM	1023	HD2	LEU	63	5.789	-6.709	3.848	1.00	0.00
ATOM	836	HA	SER	52	-1.690	-3.005	-10.027	1.00	0.00	ATOM	930	CA	THR	58	-3.409	-11.465	-0.527	1.00	0.00	ATOM	1024	HD2	LEU	63	5.789	-6.709	3.848	1.00	0.00
ATOM	837	CB	SER	52	-1.820	-1.980	-9.715	1.00	0.00	ATOM	931	HA	THR	58	-4.423	-10.177	-0.201	1.00	0.00	ATOM	1025	HD2	LEU	63	6.252	-8.049	4.565	1.00	0.00
ATOM	838	HB	SER	52	-1.447	-1.860	-10.708	1.00	0.00	ATOM	932	CB	THR	58	-4.448	-9.126	-0.112	1.00	0.00	ATOM	1026	C	LEU	63	6.204	-10.880	1.817	1.00	0.00
ATOM	839	HB	SER	52	-1.447	-1.860	-10.708	1.00	0.00	ATOM	933	HB	THR	58	-4.448	-9.126	-0.112	1.00	0.00	ATOM	1027	N	LEU	63	7.391	-11.134	2.019	1.00	0.00
ATOM	840	HB	SER	52	-0.287	-1.608	-10.877	1.00	0.00	ATOM	934	HB	THR	58	-5.974	-10.557	-1.238	1.00	0.00	ATOM	1028	HB	LEU	63	4.287	-11.235	2.565	1.00	0.00
ATOM	841	HG	SER	52	-3.993	-2.118	-8.488	1.00	0.00	ATOM	935	CO2	THR	58	-3.515	-10.598	-3.304	1.00	0.00	ATOM	1029	N	LEU	63	5.469	-12.487	3.492	1.00	0.00
ATOM	842	C	SER	52	-3.353	-2.308	-7.452	1.00	0.00	ATOM	936	CO2	THR	58	-4.236	-11.003	-3.999	1.00	0.00	ATOM	1030	CA	LVS	64	5.469	-12.487	3.492	1.00	0.00
ATOM	843	O	SER	52	-5.310	-2.379	-8.565	1.00	0.00	ATOM	937	HD2	THR	58	-2.775	-11.349	-3.073	1.00	0.00	ATOM	1031	HA	LVS	64	4.155	-13.122	3.953	1.00	0.00
ATOM	844	N	PRO	53	-6.087	-2.882	-7.435	1.00	0.00	ATOM	938	HD2	THR	58	-3.031	-9.740	-3.747	1.00	0.00	ATOM	1032	CB	LVS	64	3.506	-13.240	3.098	1.00	0.00
ATOM	845	CA	PRO	53	-6.390	-2.083	-6.773	1.00	0.00	ATOM	939	HD2	THR	58	-1.956	-9.561	-0.952	1.00	0.00	ATOM	1033	HB1	LVS	64	4.365	-14.094	4.372	1.00	0.00
ATOM	846	CA	PRO	53	-8.160	-3.415	-7.431	1.00	0.00	ATOM	940	C	THR	58	-1.095	-10.948	-0.318	1.00	0.00	ATOM	1034	HB2	LVS	64	3.316	-12.501	3.000	1.00	0.00
ATOM	847	HB1	PRO	53	-7.122	-4.519	-8.355	1.00	0.00	ATOM	941	C	THR	58	-1.095	-10.948	-0.318	1.00	0.00	ATOM	1035	HB2	LVS	64	3.316	-12.501	3.000	1.00	0.00
ATOM	848	HB2	PRO	53	-7.512	-2.661	-9.337	1.00	0.00	ATOM	942	HA	THR	58	-2.426	-8.992	-0.413	1.00	0.00	ATOM	1036	HB2	LVS	64	2.356	-12.488	4.907	1.00	0.00
ATOM	849	CO	PRO	53	-7.963	-3.263	-10.110	1.00	0.00	ATOM	943	HA	THR	58	-0.355	-8.132	-0.851	1.00	0.00	ATOM	1037	HD2	LVS	64	3.859	-12.663	6.408	1.00	0.00
ATOM	850	CO	PRO	53	-8.139	-3.182	-9.109	1.00	0.00	ATOM	944	CA	THR	58	-0.404	-6.605	-0.783	1.00	0.00	ATOM	1038	CD	LVS	64	3.007	-13.038	6.957	1.00	0.00
ATOM	851	HD1	PRO	53	-1.438	-2.187	-9.770	1.00	0.00	ATOM	945	HA	THR	58	-0.556	-6.605	-0.783	1.00	0.00	ATOM	1039	HD1	LVS	64	4.618	-11.428	6.351	1.00	0.00
ATOM	852	HD2	PRO	53	-1.438	-2.187	-9.770	1.00	0.00	ATOM	946	CB	THR	58	-1.260	-6.605	-1.112	1.00	0.00	ATOM	1040	HD2	LVS	64	4.426	-11.459	7.141	1.00	0.00
ATOM	853	CD	PRO	53	-1.438	-2.187	-9.770	1.00	0.00	ATOM	947	HB2	THR	58	-1.260	-6.605	-1.112	1.00	0.00	ATOM	1041	CE	LVS	64	4.711	-10.712	5.614	1.00	0.00
ATOM	854	HB1	PRO	53	-8.160	-3.415	-7.431	1.00	0.00	ATOM	948	HB2	THR	58	-1.260	-6.605	-1.112	1.00	0.00	ATOM	1042	HB3	LVS	64	4.711	-10.712	5.614	1.00	0.00
ATOM	855	HB2	PRO	53	-8.160	-3.415	-7.431	1.00	0.00	ATOM	949	HB1	THR	58	-1.503	-6.385	1.032	1.00	0.00	ATOM	1043	HB3	LVS	64	4.711	-10.712	5.614	1.00	0.00
ATOM	856	C	PRO	53	-5.340	-3.947	-6.943	1.00	0.00	ATOM	950	HOL	THR	58	-0.218	-6.385	1.032	1.00	0.00	ATOM	1044	HB2	LVS	64	6.086	-12.655	7.569	1.00	0.00
ATOM	857	O	PRO	53	-5.425	-5.137	-6.943	1.00	0.00	ATOM	951	HOL	THR	58	-0.218	-6.385	1.032	1.00	0.00	ATOM	1045	HB2	LVS	64	6.086	-12.655	7.569	1.00	0.00
ATOM	858	N	MET	54	-4.609	-3.512	-5.619	1.00	0.00	ATOM	952	CE	THR	58	-0.969	-4.397	0.643	1.00	0.00	ATOM	1046	HB2	LVS	64	6.295	-11.028	7.968	1.00	0.00
ATOM	859	N	MET	54	-4.591	-3.512	-5.619	1.00	0.00	ATOM	953	CE	THR	58	-0.969	-4.397	0.643	1.00	0.00	ATOM	1047	HB2	LVS	64	6.295	-11.028	7.968	1.00	0.00
ATOM	860	CA	MET	54	-8.950	-4.428	-4.777	1.00	0.00	ATOM	954	HB1	THR	58	1.178	-4.324	-1.443	1.00	0.00	ATOM	1048	C	LVS	64	6.405	-14.062	3.973	1.00	0.00
ATOM	861	CA	MET	54	-8.950	-4.428	-4.777	1.00	0.00	ATOM	955	HB2	THR	58	1.178	-4.324	-1.443	1.00	0.00	ATOM	1049	C	LVS	64	6.405	-14.062	3.973	1.00	0.00
ATOM	862	CB	MET	54	-2.398	-4.508	-4.253	1.00	0.00	ATOM	956	HB2	THR	58	1.178	-4.324	-1.443	1.00	0.00	ATOM	1050	C	LVS	64	6.405	-14.062	3.973	1.00	0.00
ATOM	863	HB1	MET	54	-2.390	-4.624	-6.327	1.00	0.00	ATOM	957	C	THR	58	0.582	-6.669	0.228	1.00	0.00	ATOM	1051	HN	ASN	65	5.569	-13.420	1.043	1.00	0.00
ATOM	864	HB1	MET	54	-1.897	-3.587	-4.995	1.00	0.00	ATOM	958	O	THR	58	1.701	-9.092	-0.061	1.00	0.00	ATOM	1052	CA	ASN	65	7.103	-14.872	1.023	1.00	0.00
ATOM	865	CG	MET	54	-1.615	-5.664	-4.644	1.00	0.00	ATOM	959	N	SER	60	0.126	-8.821	1.476	1.00	0.00	ATOM	1053	HA	ASN	65	7.632	-15.410	1.795	1.00	0.00
ATOM	866	HOL	MET	54	-1.672	-5.664	-4.592	1.00	0.00	ATOM	960	N	SER	60	-0.772	-8.821	1.644	1.00	0.00	ATOM	1054	CA	ASN	65	6.254	-15.855	0.242	1.00	0.00
ATOM	867	HOL	MET	54	-6.653	-5.720	-5.132	1.00	0.00	ATOM	961	CA	SER	60	0.925	-9.088	2.601	1.00	0.00	ATOM	1055	HB1	ASN	65	6.762	-16.082	-0.712	1.00	0.00
ATOM																													

REPLACEMENT SHEET

ATOM	1101	HE1	THR	67	6.089	-16.237	-6.940	1.00	0.00	ATOM	1195	NE1	LVS	72	7.488	3.345	-12.403	1.00	0.00	ATOM	1389	HD1	LEU	78	0.056	-1.174	1.350	1.00	0.00
ATOM	1102	CE2	THR	67	4.788	-13.123	-7.044	1.00	0.00	ATOM	1196	HE1	LVS	72	7.488	3.345	-12.403	1.00	0.00	ATOM	1390	HD2	LEU	78	0.251	-1.375	1.350	1.00	0.00
ATOM	1103	HE2	THR	67	4.281	-12.449	-7.711	1.00	0.00	ATOM	1197	HE2	LVS	72	5.902	3.166	-13.386	1.00	0.00	ATOM	1391	HD3	LEU	78	1.327	-1.855	0.521	1.00	0.00
ATOM	1104	CE	THR	67	5.153	-14.337	-7.466	1.00	0.00	ATOM	1198	HE3	LVS	72	6.961	2.316	-13.037	1.00	0.00	ATOM	1392	CD1	LEU	78	1.094	0.844	2.605	1.00	0.00
ATOM	1105	CH	THR	67	4.850	-14.793	-8.745	1.00	0.00	ATOM	1199	C	LVS	72	5.915	1.152	-7.680	1.00	0.00	ATOM	1393	HD21	LEU	78	1.126	1.080	2.303	1.00	0.00
ATOM	1106	C	THR	67	4.028	-15.286	-8.737	1.00	0.00	ATOM	1200	C	LVS	72	4.723	-0.245	-7.305	1.00	0.00	ATOM	1394	HD22	LEU	78	1.641	0.721	3.529	1.00	0.00
ATOM	1107	CH	THR	67	4.850	-15.286	-8.737	1.00	0.00	ATOM	1201	N	LEU	73	6.347	-2.445	-7.305	1.00	0.00	ATOM	1395	HD23	LEU	78	0.068	0.069	-1.009	1.00	0.00
ATOM	1108	C	THR	67	6.950	-10.676	-3.063	1.00	0.00	ATOM	1202	N	LEU	73	5.446	-2.445	-7.305	1.00	0.00	ATOM	1396	CD1	LEU	78	0.780	2.760	-1.231	1.00	0.00
ATOM	1109	N	THR	68	6.859	-9.937	-2.995	1.00	0.00	ATOM	1203	CA	LEU	73	4.802	-3.527	-7.966	1.00	0.00	ATOM	1397	HA	THR	83	-4.515	6.449	-2.129	1.00	0.00
ATOM	1110	HN	THR	68	6.909	-10.187	-1.978	1.00	0.00	ATOM	1204	HA	LEU	73	6.231	-4.764	-6.964	1.00	0.00	ATOM	1398	HN	GLN	79	1.649	2.393	-2.191	1.00	0.00
ATOM	1111	CA	THR	68	6.515	-8.568	-3.280	1.00	0.00	ATOM	1205	CB	LEU	73	7.432	-4.050	-5.922	1.00	0.00	ATOM	1399	HN	GLN	79	0.410	4.097	-2.184	1.00	0.00
ATOM	1112	HA	THR	68	6.570	-8.478	-4.335	1.00	0.00	ATOM	1206	HB1	LEU	73	6.150	-4.656	-7.142	1.00	0.00	ATOM	1300	HA	GLN	79	0.357	4.744	-1.221	1.00	0.00
ATOM	1113	HN	THR	68	5.089	-8.253	-8.821	1.00	0.00	ATOM	1207	HB2	LEU	73	5.308	-4.614	-7.466	1.00	0.00	ATOM	1301	HA	GLN	79	1.478	4.631	-3.154	1.00	0.00
ATOM	1114	HN	THR	68	5.089	-8.253	-8.821	1.00	0.00	ATOM	1208	CB	LEU	73	5.308	-4.614	-7.466	1.00	0.00	ATOM	1302	CB	GLN	79	1.727	3.631	-3.850	1.00	0.00
ATOM	1115	HB2	THR	68	5.089	-8.253	-8.821	1.00	0.00	ATOM	1209	CD1	LEU	73	6.313	-7.925	-3.140	1.00	0.00	ATOM	1303	HB1	GLN	79	1.052	5.860	-2.563	1.00	0.00
ATOM	1116	CB	THR	68	4.089	-8.103	-3.946	1.00	0.00	ATOM	1210	CD1	LEU	73	6.313	-7.925	-3.140	1.00	0.00	ATOM	1304	HB2	GLN	79	0.674	5.507	-4.913	1.00	0.00
ATOM	1117	CD1	THR	68	4.278	-8.079	-5.174	1.00	0.00	ATOM	1211	HD1	LEU	73	7.328	-7.136	-7.406	1.00	0.00	ATOM	1305	HG1	GLN	79	0.792	6.356	-3.421	1.00	0.00
ATOM	1118	HD1	THR	68	5.157	-9.336	-5.327	1.00	0.00	ATOM	1212	HD2	LEU	73	5.868	-8.115	-6.961	1.00	0.00	ATOM	1306	HG2	GLN	79	2.194	6.807	-4.202	1.00	0.00
ATOM	1119	CD2	THR	68	2.946	-7.334	-1.772	1.00	0.00	ATOM	1213	HD3	LEU	73	5.247	-5.193	-6.961	1.00	0.00	ATOM	1307	HG3	GLN	79	2.522	7.551	-3.307	1.00	0.00
ATOM	1120	HD2	THR	68	2.783	-6.844	-2.823	1.00	0.00	ATOM	1214	CD2	LEU	73	4.982	-6.889	-9.345	1.00	0.00	ATOM	1308	OE1	GLN	79	2.766	7.001	-3.420	1.00	0.00
ATOM	1121	HE1	THR	68	3.523	-8.352	-6.136	1.00	0.00	ATOM	1215	HD21	LEU	73	4.439	-5.552	-9.345	1.00	0.00	ATOM	1309	OE2	GLN	79	2.996	6.807	-3.420	1.00	0.00
ATOM	1122	HE2	THR	68	3.523	-8.352	-6.136	1.00	0.00	ATOM	1216	HD22	LEU	73	4.439	-5.552	-9.345	1.00	0.00	ATOM	1310	NE2	GLN	79	2.522	7.551	-3.307	1.00	0.00
ATOM	1123	CE2	THR	68	2.022	-7.185	-4.789	1.00	0.00	ATOM	1217	HD23	LEU	73	4.587	-3.247	-5.881	1.00	0.00	ATOM	1311	NE3	GLN	79	2.350	6.182	-2.083	1.00	0.00
ATOM	1124	HE2	THR	68	1.141	-6.581	-4.633	1.00	0.00	ATOM	1218	C	LEU	73	3.375	-3.053	-5.958	1.00	0.00	ATOM	1312	HE22	GLN	79	0.953	4.061	-2.970	1.00	0.00
ATOM	1125	CE	THR	68	2.233	-7.811	-5.998	1.00	0.00	ATOM	1219	C	LEU	73	5.218	-3.316	-4.633	1.00	0.00	ATOM	1313	C	GLN	79	-0.953	4.061	-2.970	1.00	0.00
ATOM	1126	CH	THR	68	1.315	-7.655	-7.032	1.00	0.00	ATOM	1220	N	PHIE	74	6.184	-3.474	-4.633	1.00	0.00	ATOM	1314	N	GLN	79	-1.745	4.997	-2.768	1.00	0.00
ATOM	1127	HN	THR	68	0.444	-7.518	-6.637	1.00	0.00	ATOM	1221	HN	PHIE	74	4.223	-6.043	-1.166	1.00	0.00	ATOM	1315	N	GLN	79	-1.205	2.961	-3.663	1.00	0.00
ATOM	1128	C	THR	68	7.484	-7.510	-2.656	1.00	0.00	ATOM	1222	CA	PHIE	74	4.223	-6.043	-1.166	1.00	0.00	ATOM	1316	HN	ARG	80	-0.522	2.263	-3.663	1.00	0.00
ATOM	1129	C	THR	68	7.484	-7.510	-2.656	1.00	0.00	ATOM	1223	CA	PHIE	74	2.589	-2.622	-3.526	1.00	0.00	ATOM	1317	CA	ARG	80	-2.439	3.755	-4.287	1.00	0.00
ATOM	1130	N	VAL	69	8.744	-7.567	-2.529	1.00	0.00	ATOM	1224	HN	PHIE	74	6.174	-3.924	-2.095	1.00	0.00	ATOM	1318	CA	ARG	80	-1.151	1.505	-5.449	1.00	0.00
ATOM	1131	HN	VAL	69	9.000	-8.882	-2.895	1.00	0.00	ATOM	1225	HB1	PHIE	74	5.587	-4.799	-2.614	1.00	0.00	ATOM	1319	HB1	ARG	80	-2.686	0.787	-5.959	1.00	0.00
ATOM	1132	CA	VAL	69	9.761	-7.086	-1.962	1.00	0.00	ATOM	1226	HB2	PHIE	74	4.504	-3.929	-1.033	1.00	0.00	ATOM	1320	CG	ARG	80	-2.053	2.063	-6.740	1.00	0.00
ATOM	1133	CA	VAL	69	9.284	-6.461	-1.221	1.00	0.00	ATOM	1227	CD1	PHIE	74	4.221	-6.043	-1.166	1.00	0.00	ATOM	1321	HG1	ARG	80	-2.053	2.063	-6.740	1.00	0.00
ATOM	1134	CB	VAL	69	10.907	-7.854	-1.276	1.00	0.00	ATOM	1228	CD1	PHIE	74	4.223	-6.043	-1.166	1.00	0.00	ATOM	1322	HG2	ARG	80	-2.742	1.180	-7.363	1.00	0.00
ATOM	1135	CB	VAL	69	10.713	-7.954	-1.984	1.00	0.00	ATOM	1229	HD1	PHIE	74	4.223	-6.043	-1.166	1.00	0.00	ATOM	1323	HG3	ARG	80	-4.122	2.649	-6.882	1.00	0.00
ATOM	1136	CD1	VAL	69	9.734	-4.165	-4.164	1.00	0.00	ATOM	1230	CD2	PHIE	74	3.554	-4.232	2.257	1.00	0.00	ATOM	1324	HD1	ARG	80	-4.122	2.649	-6.882	1.00	0.00
ATOM	1137	HD1	VAL	69	10.352	-6.139	-3.043	1.00	0.00	ATOM	1231	CD2	PHIE	74	3.554	-4.232	2.257	1.00	0.00	ATOM	1325	HD2	ARG	80	-4.782	1.982	-7.393	1.00	0.00
ATOM	1138	HD2	VAL	69	9.487	-9.183	-0.393	1.00	0.00	ATOM	1232	CE1	PHIE	74	3.006	-6.238	0.930	1.00	0.00	ATOM	1326	HD3	ARG	80	-4.186	3.966	-7.310	1.00	0.00
ATOM	1139	HD3	VAL	69	11.171	-9.640	-0.133	1.00	0.00	ATOM	1233	CE1	PHIE	74	3.006	-6.238	0.930	1.00	0.00	ATOM	1327	HD4	ARG	80	-4.186	3.966	-7.310	1.00	0.00
ATOM	1140	CG2	VAL	69	10.423	-7.059	-0.990	1.00	0.00	ATOM	1234	CE2	PHIE	74	3.518	-2.929	1.063	1.00	0.00	ATOM	1328	NE	ARG	80	-4.186	3.966	-7.310	1.00	0.00
ATOM	1141	CG2	VAL	69	11.422	-7.059	-0.990	1.00	0.00	ATOM	1235	CE2	PHIE	74	3.518	-2.929	1.063	1.00	0.00	ATOM	1329	HE	ARG	80	-4.186	3.966	-7.310	1.00	0.00
ATOM	1142	HE22	VAL	69	12.197	-7.619	-0.411	1.00	0.00	ATOM	1236	CE3	PHIE	74	2.584	-4.232	2.257	1.00	0.00	ATOM	1330	HE	ARG	80	-4.186	3.966	-7.310	1.00	0.00
ATOM	1143	HE23	VAL	69	10.608	-6.872	-0.597	1.00	0.00	ATOM	1237	CE3	PHIE	74	2.584	-4.232	2.257	1.00	0.00	ATOM	1331	NH1	ARG	80	-6.486	4.030	-7.305	1.00	0.00
ATOM	1144	C	VAL	69	10.352	-6.139	-3.043	1.00	0.00	ATOM	1238	CE3	PHIE	74	2.584	-4.232	2.257	1.00	0.00	ATOM	1332	NH2	ARG	80	-4.955	4.505	-6.213	1.00	0.00
ATOM	1145	C	VAL	69	11.543	-6.272	-3.345	1.00	0.00	ATOM	1239	C	PHIE	74	3.172	-1.236	-2.879	1.00	0.00	ATOM	1333	NH3	ARG	80	-5.294	5.750	-6.168	1.00	0.00
ATOM	1146	N	SER	70	9.507	-5.372	-3.634	1.00	0.00	ATOM	1240	N	PHIE	75	5.392	-3.092	-3.095	1.00	0.00	ATOM	1334	NH2	ARG	80	-6.418	6.219	-8.386	1.00	0.00
ATOM	1147	HN	SER	70	8.566	-5.371	-3.359	1.00	0.00	ATOM	1241	CA	PHIE	75	6.262	-1.309	-3.281	1.00	0.00	ATOM	1335	NH21	ARG	80	-6.418	6.219	-8.386	1.00	0.00
ATOM	1148	CA	SER	70	9.934	-4.431	-4.003	1.00	0.00	ATOM	1242	CA	PHIE	75	5.392	-3.092	-3.281	1.00	0.00	ATOM	1336	NH22	ARG	80	-6.418	6.219	-8.386	1.00	0.00
ATOM	1149	CB	SER	70	9.734	-4.165	-4.164	1.00	0.00	ATOM	1243	CA	PHIE</																

REPLACEMENT SHEET

ATOM	1383	CE2 THR	83	-4.464	7.286	-5.619	1.00	0.00	ATOM	1477	H81 ASN	89	-14.820	8.581	-0.042	1.00	0.00	ATOM	1571	O	TYR	95	-10.170	6.352	2.951	1.00	0.00		
ATOM	1384	THR	83	-4.188	7.286	-5.619	1.00	0.00	ATOM	1478	H82 ASN	89	-13.706	7.286	-1.173	1.00	0.00	ATOM	1572	N	TYR	96	-11.743	7.934	3.443	1.00	0.00		
ATOM	1385	H822 THR	83	-4.884	8.255	-6.049	1.00	0.00	ATOM	1479	H83 ASN	89	-12.001	7.798	-0.149	1.00	0.00	ATOM	1573	HN	TYR	96	-10.862	7.934	3.443	1.00	0.00		
ATOM	1386	THR	83	-5.228	6.656	-6.049	1.00	0.00	ATOM	1480	CO1 ASN	89	-12.001	7.798	-0.149	1.00	0.00	ATOM	1574	CA	TYR	96	-10.862	7.934	3.443	1.00	0.00		
ATOM	1387	O	THR	83	-5.278	6.017	-3.711	1.00	0.00	ATOM	1481	H81 ASN	89	-13.444	6.136	-0.609	1.00	0.00	ATOM	1575	HA	TYR	96	-11.579	8.795	2.268	1.00	0.00	
ATOM	1388	O	THR	83	-7.221	6.740	-4.033	1.00	0.00	ATOM	1482	H82 ASN	89	-12.769	5.467	-0.377	1.00	0.00	ATOM	1576	CB	TYR	96	-11.579	10.321	2.712	1.00	0.00	
ATOM	1389	N	ASN	84	-5.333	4.690	-3.756	1.00	0.00	ATOM	1483	H822 ASN	89	-14.348	5.910	-0.952	1.00	0.00	ATOM	1577	H81 TYR	96	-12.376	10.073	2.027	1.00	0.00		
ATOM	1390	HN	ASN	84	-5.350	4.170	-3.476	1.00	0.00	ATOM	1484	O	ASN	89	-15.944	5.048	-0.252	1.00	0.00	ATOM	1578	H82 TYR	96	-10.717	11.579	6.460	1.00	0.00	
ATOM	1391	HN	ASN	84	-5.350	4.170	-3.476	1.00	0.00	ATOM	1485	N	ASN	89	-15.701	10.222	-2.272	1.00	0.00	ATOM	1579	CO	TYR	96	-10.717	11.579	6.460	1.00	0.00
ATOM	1392	CA	ASN	84	-7.822	4.170	-3.476	1.00	0.00	ATOM	1486	N	ASN	89	-15.701	10.222	-2.272	1.00	0.00	ATOM	1580	H81 TYR	96	-10.717	11.579	6.460	1.00	0.00	
ATOM	1393	CB	ASN	84	-7.250	2.481	-4.321	1.00	0.00	ATOM	1487	HN	ALA	90	-17.141	7.728	-3.446	1.00	0.00	ATOM	1581	H81 TYR	96	-10.717	11.579	6.460	1.00	0.00	
ATOM	1394	H81 ASN	84	-6.457	2.342	-5.039	1.00	0.00	ATOM	1488	CA	ALA	90	-17.995	6.655	-3.783	1.00	0.00	ATOM	1582	CD2 TYR	96	-10.364	12.653	3.758	1.00	0.00		
ATOM	1395	H82 ASN	84	-6.939	2.107	-3.362	1.00	0.00	ATOM	1489	HA	ALA	90	-17.468	10.184	-4.599	1.00	0.00	ATOM	1583	H82 TYR	96	-10.714	12.653	3.758	1.00	0.00		
ATOM	1396	CO	ASN	84	-6.464	1.707	-4.778	1.00	0.00	ATOM	1490	H81 ALA	90	-19.185	8.946	-4.336	1.00	0.00	ATOM	1584	CE1 TYR	96	-9.471	12.183	0.159	1.00	0.00		
ATOM	1397	CO1 ASN	84	-6.459	2.282	-5.218	1.00	0.00	ATOM	1491	H82 ALA	90	-19.230	7.944	-3.946	1.00	0.00	ATOM	1585	H81 TYR	96	-9.120	12.073	-0.855	1.00	0.00			
ATOM	1398	H822 ASN	84	-7.564	0.405	-4.512	1.00	0.00	ATOM	1492	H82 ALA	90	-19.128	8.150	-3.404	1.00	0.00	ATOM	1586	CE2 TYR	96	-9.376	13.501	2.155	1.00	0.00			
ATOM	1399	H822 ASN	84	-7.564	0.405	-4.512	1.00	0.00	ATOM	1493	O	ALA	90	-18.356	10.662	-2.705	1.00	0.00	ATOM	1587	HN	TYR	96	-10.352	8.608	5.478	1.00	0.00	
ATOM	1400	H822 ASN	84	-6.158	-0.147	-4.960	1.00	0.00	ATOM	1494	O	ALA	90	-18.356	10.662	-2.705	1.00	0.00	ATOM	1588	CE2 TYR	96	-8.347	14.280	0.265	1.00	0.00		
ATOM	1401	O	ASN	84	-6.670	4.212	-3.201	1.00	0.00	ATOM	1495	O	ALA	90	-18.356	10.662	-2.705	1.00	0.00	ATOM	1589	CH TYR	96	-8.347	14.280	0.265	1.00	0.00	
ATOM	1402	C	ASN	84	-6.956	4.879	-3.514	1.00	0.00	ATOM	1496	N	PRO	91	-18.657	10.179	-1.487	1.00	0.00	ATOM	1590	HN TYR	96	-8.832	14.692	-0.455	1.00	0.00	
ATOM	1403	N	CYS	85	-6.520	3.657	-2.003	1.00	0.00	ATOM	1497	CA	PRO	91	-19.087	11.013	-0.781	1.00	0.00	ATOM	1591	O	TYR	96	-10.009	9.267	4.346	1.00	0.00
ATOM	1404	HN	CYS	85	-7.708	3.142	-1.812	1.00	0.00	ATOM	1498	HA	PRO	91	-19.505	11.951	-0.717	1.00	0.00	ATOM	1592	C	TYR	96	-8.884	9.764	4.303	1.00	0.00
ATOM	1405	HN	CYS	85	-7.708	3.142	-1.812	1.00	0.00	ATOM	1499	HN	PRO	91	-20.276	10.349	-1.389	1.00	0.00	ATOM	1593	HN	TYR	97	-10.532	8.608	5.478	1.00	0.00
ATOM	1406	CA	CYS	85	-10.414	3.205	-0.354	1.00	0.00	ATOM	1500	HN1 PRO	91	-21.115	10.323	-0.263	1.00	0.00	ATOM	1594	HN	TYR	97	-9.937	9.050	6.771	1.00	0.00	
ATOM	1407	CB	CYS	85	-9.028	3.205	-0.354	1.00	0.00	ATOM	1501	H82 PRO	91	-19.704	8.730	-0.022	1.00	0.00	ATOM	1595	CA	CYS	97	-9.937	9.050	6.771	1.00	0.00	
ATOM	1408	H81 CYS	85	-9.789	3.329	1.109	1.00	0.00	ATOM	1502	CO	PRO	91	-20.537	8.104	-0.288	1.00	0.00	ATOM	1596	HA	LVS	97	-10.965	8.901	7.899	1.00	0.00	
ATOM	1409	H82 CYS	85	-8.132	3.724	0.665	1.00	0.00	ATOM	1503	H81 PRO	91	-19.249	8.358	-1.094	1.00	0.00	ATOM	1597	CB	LVS	97	-11.380	7.906	7.899	1.00	0.00		
ATOM	1410	SG	CYS	85	-8.635	1.443	0.357	1.00	0.00	ATOM	1504	CO2 PRO	91	-18.679	8.764	-1.094	1.00	0.00	ATOM	1598	H81 LVS	97	-11.457	9.033	8.845	1.00	0.00		
ATOM	1411	HG	CYS	85	-9.362	0.955	0.652	1.00	0.00	ATOM	1505	CO	PRO	91	-17.943	8.153	-0.744	1.00	0.00	ATOM	1599	H82 LVS	97	-12.100	9.903	7.150	1.00	0.00	
ATOM	1412	HG	CYS	85	-10.966	5.413	-0.398	1.00	0.00	ATOM	1506	H82 PRO	91	-17.943	8.153	-0.744	1.00	0.00	ATOM	1600	H81 LVS	97	-12.100	9.903	7.150	1.00	0.00		
ATOM	1413	C	CYS	85	-10.966	5.413	-0.398	1.00	0.00	ATOM	1507	HN1 PRO	91	-17.943	8.153	-0.744	1.00	0.00	ATOM	1601	H82 LVS	97	-12.100	9.903	7.150	1.00	0.00		
ATOM	1414	N	LVS	86	-8.898	6.150	-0.993	1.00	0.00	ATOM	1508	C	PRO	91	-17.976	11.268	0.633	1.00	0.00	ATOM	1602	H82 LVS	97	-11.711	10.849	10.972	1.00	0.00	
ATOM	1415	HN	LVS	86	-8.021	5.820	-1.381	1.00	0.00	ATOM	1509	O	PRO	91	-18.242	11.671	1.765	1.00	0.00	ATOM	1603	CA	LVS	97	-13.665	10.674	9.042	1.00	0.00
ATOM	1416	CA	LVS	86	-9.102	7.586	-0.940	1.00	0.00	ATOM	1510	N	GLU	92	-16.734	11.671	0.234	1.00	0.00	ATOM	1604	H81 LVS	97	-12.990	9.142	9.601	1.00	0.00	
ATOM	1417	HA	LVS	86	-9.725	7.741	0.029	1.00	0.00	ATOM	1511	HN	GLU	92	-16.582	10.671	1.132	1.00	0.00	ATOM	1605	CE	LVS	97	-11.842	10.857	9.131	1.00	0.00
ATOM	1418	CB	LVS	86	-7.758	8.280	-0.623	1.00	0.00	ATOM	1512	CA	GLU	92	-15.956	11.174	1.132	1.00	0.00	ATOM	1606	CE	LVS	97	-11.842	10.857	9.131	1.00	0.00
ATOM	1419	HN	LVS	86	-7.078	7.980	-1.405	1.00	0.00	ATOM	1513	HN	GLU	92	-15.956	11.174	1.132	1.00	0.00	ATOM	1607	HN	LVS	97	-11.842	10.857	9.131	1.00	0.00
ATOM	1420	H82 LVS	86	-7.853	9.807	-0.532	1.00	0.00	ATOM	1514	HN1 GLU	92	-15.140	12.750	1.408	1.00	0.00	ATOM	1608	H82 LVS	97	-12.411	12.111	10.629	1.00	0.00			
ATOM	1421	CG	LVS	86	-8.858	10.019	-0.901	1.00	0.00	ATOM	1515	H81 GLU	92	-16.143	13.150	2.000	1.00	0.00	ATOM	1609	HN	LVS	97	-12.411	12.111	10.629	1.00	0.00	
ATOM	1422	H81 LVS	86	-8.858	10.019	-0.901	1.00	0.00	ATOM	1516	H82 GLU	92	-16.143	13.150	2.000	1.00	0.00	ATOM	1610	HN	LVS	97	-12.411	12.111	10.629	1.00	0.00		
ATOM	1423	H82 LVS	86	-7.625	10.178	0.356	1.00	0.00	ATOM	1517	CO	GLU	92	-15.139	13.487	0.150	1.00	0.00	ATOM	1611	H82 LVS	97	-12.672	12.013	11.666	1.00	0.00		
ATOM	1424	CA	LVS	86	-6.882	10.454	-1.626	1.00	0.00	ATOM	1518	H81 GLU	92	-15.963	14.133	0.021	1.00	0.00	ATOM	1612	H23 LVS	97	-11.362	12.289	10.123	1.00	0.00		
ATOM	1425	H81 LVS	86	-4.931	11.050	-2.334	1.00	0.00	ATOM	1519	CO2 GLU	92	-15.963	14.133	0.021	1.00	0.00	ATOM	1613	C	LVS	97	-8.816	8.031	6.955	1.00	0.00		
ATOM	1426	HN	LVS	86	-5.860	11.308	-0.930	1.00	0.00	ATOM	1520	CO1 GLU	92	-13.118	14.335	1.276	1.00	0.00	ATOM	1614	HN	CYS	98	-6.725	6.681	1.00	0.00		
ATOM	1427	CB	LVS	86	-5.744	10.970	-0.990	1.00	0.00	ATOM	1521	CO1 GLU	92	-13.118	14.335	1.276	1.00	0.00	ATOM	1615	N	CYS	98	-10.105	6.559	6.681	1.00	0.00	
ATOM	1428	H81 LVS	86	-6.916	11.231	-1.447	1.00	0.00	ATOM	1522	CO2 GLU	92	-13.679	15.021	1.276	1.00	0.00	ATOM	1616	HN	CYS	98	-8.203	5.671	7.039	1.00	0.00		
ATOM	1429	H82 LVS	86	-6.276	12.778	-0.922	1.00	0.00	ATOM	1523	C	GLU	92	-15.843	10.432	2.436	1.00	0.00	ATOM	1617	CA	CYS	98	-8.203	5.671	7.039	1.00	0.00	
ATOM	1430	NZ	LVS	86	-7.207	13.135	-0.941	1.00	0.00	ATOM	1524	O	GLU	92	-15.529	10.925	3.519	1.00	0.00	ATOM	1618	HA	CYS	98	-8.816	4.348	6.588	1.00	0.00
ATOM	1431	H82 LVS	86	-7.207	13.135	-0.941	1.00	0.00	ATOM	1525	N	SER	93	-16.427	9.248	2.313	1.00	0.00	ATOM	1619	CB	CYS	98	-8.816	4.348	6.588	1.00	0.00	
ATOM	1432	H82 LVS	86	-7.260	12.855	-1.448	1.00	0.00	ATOM	1526	HN																		

REPLACEMENT SHEET

ATOM	1665	HE2	TLE	101	-6.705	5.862	11.709	1.00	0.00	ATOM	1759	CE2	PHE	106	3.941	2.500	1.491	1.00	0.00	ATOM	1853	NZ	LVS	111	7.860	11.169	2.702	1.00	0.00
ATOM	1666	HE3	TLE	101	-7.384	5.966	10.534	1.00	0.00	ATOM	1760	HE2	PHE	106	3.995	2.653	0.471	1.00	0.00	ATOM	1854	HE2	LVS	111	7.860	11.560	3.543	1.00	0.00
ATOM	1667	C	TLE	101	-2.511	6.457	8.326	1.00	0.00	ATOM	1761	CE	PHE	106	4.605	4.780	1.854	1.00	0.00	ATOM	1855	HE2	LVS	111	7.151	10.571	2.187	1.00	0.00
ATOM	1668	N	LVS	102	-1.335	6.807	8.332	1.00	0.00	ATOM	1762	HE2	PHE	106	5.125	5.734	1.050	1.00	0.00	ATOM	1856	HE2	LVS	111	8.121	11.954	2.084	1.00	0.00
ATOM	1669	N	LVS	102	-2.876	5.343	7.590	1.00	0.00	ATOM	1763	C	PHE	106	4.388	5.073	5.110	1.00	0.00	ATOM	1857	C	LVS	111	12.448	7.070	4.423	1.00	0.00
ATOM	1670	N	LVS	102	-2.829	5.117	7.527	1.00	0.00	ATOM	1764	C	PHE	106	4.388	5.073	5.110	1.00	0.00	ATOM	1858	C	LVS	111	12.448	7.070	4.423	1.00	0.00
ATOM	1671	CA	LVS	102	-1.350	4.851	6.972	1.00	0.00	ATOM	1765	N	PHE	107	3.795	6.155	4.434	1.00	0.00	ATOM	1859	N	LVS	112	12.346	6.255	5.421	1.00	0.00
ATOM	1672	CA	LVS	102	-1.350	4.851	6.972	1.00	0.00	ATOM	1766	CA	PHE	107	4.575	7.140	3.686	1.00	0.00	ATOM	1860	CA	LVS	112	13.485	5.516	5.983	1.00	0.00
ATOM	1673	CA	LVS	102	-2.569	3.416	6.082	1.00	0.00	ATOM	1767	CA	PHE	107	5.076	6.516	2.885	1.00	0.00	ATOM	1861	CA	LVS	112	13.485	5.516	5.983	1.00	0.00
ATOM	1674	HE1	LVS	102	-3.630	3.443	6.283	1.00	0.00	ATOM	1768	HA	PHE	107	3.662	8.212	3.077	1.00	0.00	ATOM	1862	HA	LVS	112	14.297	6.205	6.151	1.00	0.00
ATOM	1675	HE2	LVS	102	-2.408	3.700	5.053	1.00	0.00	ATOM	1769	HA	PHE	107	3.662	8.212	3.077	1.00	0.00	ATOM	1863	CA	LVS	112	13.126	4.819	7.296	1.00	0.00
ATOM	1676	CA	LVS	102	-2.083	1.971	6.260	1.00	0.00	ATOM	1770	HE1	PHE	107	4.263	9.070	2.810	1.00	0.00	ATOM	1864	HE2	LVS	112	12.159	4.350	7.189	1.00	0.00
ATOM	1677	CA	LVS	102	-2.423	1.382	5.440	1.00	0.00	ATOM	1771	HE2	PHE	107	4.263	9.070	2.810	1.00	0.00	ATOM	1865	HE2	LVS	112	13.864	4.058	7.506	1.00	0.00
ATOM	1678	HE2	LVS	102	-2.423	1.382	5.440	1.00	0.00	ATOM	1772	CE	PHE	107	3.918	6.170	1.938	1.00	0.00	ATOM	1866	CA	LVS	112	13.070	5.760	9.468	1.00	0.00
ATOM	1679	HE2	LVS	102	-1.945	1.364	7.552	1.00	0.00	ATOM	1773	CE	PHE	107	3.918	6.170	1.938	1.00	0.00	ATOM	1867	CA	LVS	112	13.070	5.760	9.468	1.00	0.00
ATOM	1680	HE2	LVS	102	-2.958	0.338	7.332	1.00	0.00	ATOM	1774	HE1	PHE	107	4.674	5.771	1.863	1.00	0.00	ATOM	1868	HE2	LVS	112	12.644	6.700	8.165	1.00	0.00
ATOM	1681	HE2	LVS	102	-3.549	1.928	7.818	1.00	0.00	ATOM	1775	CE2	PHE	107	2.068	8.667	1.560	1.00	0.00	ATOM	1869	CA	LVS	112	14.842	5.027	9.087	1.00	0.00
ATOM	1682	CE2	LVS	102	-0.559	1.903	6.290	1.00	0.00	ATOM	1776	HE2	PHE	107	1.995	9.670	1.160	1.00	0.00	ATOM	1870	CE2	LVS	112	14.842	5.027	9.087	1.00	0.00
ATOM	1683	HE2	LVS	102	-0.147	2.768	5.827	1.00	0.00	ATOM	1777	CE1	PHE	107	2.356	6.069	0.214	1.00	0.00	ATOM	1871	CE2	LVS	112	15.103	6.984	8.641	1.00	0.00
ATOM	1684	HE2	LVS	102	-0.284	1.057	5.749	1.00	0.00	ATOM	1778	HE1	PHE	107	2.460	5.069	-0.154	1.00	0.00	ATOM	1872	C	LVS	112	15.103	6.984	8.641	1.00	0.00
ATOM	1685	HE2	LVS	102	-0.280	1.861	7.113	1.00	0.00	ATOM	1779	CE2	PHE	107	1.413	8.268	0.025	1.00	0.00	ATOM	1873	C	LVS	112	15.128	4.228	4.799	1.00	0.00
ATOM	1686	HE2	LVS	102	-1.945	1.364	7.552	1.00	0.00	ATOM	1780	CE2	PHE	107	1.413	8.268	0.025	1.00	0.00	ATOM	1874	CA	LVS	114	13.760	5.460	9.454	1.00	0.00
ATOM	1687	C	LVS	102	-2.220	7.612	3.122	1.00	0.00	ATOM	1781	CE	PHE	107	1.542	6.977	-0.449	1.00	0.00	ATOM	1875	CA	LVS	114	13.760	5.460	9.454	1.00	0.00
ATOM	1688	N	LVS	103	-1.414	6.265	5.433	1.00	0.00	ATOM	1782	CE	PHE	107	1.542	6.977	-0.449	1.00	0.00	ATOM	1876	CA	LVS	114	13.760	5.460	9.454	1.00	0.00
ATOM	1689	N	LVS	103	-2.377	6.439	5.484	1.00	0.00	ATOM	1783	C	PHE	107	5.636	7.787	4.571	1.00	0.00	ATOM	1877	CA	LVS	114	14.151	2.356	3.550	1.00	0.00
ATOM	1690	CA	LVS	103	-0.581	7.132	4.612	1.00	0.00	ATOM	1784	C	PHE	107	5.636	7.787	4.571	1.00	0.00	ATOM	1878	CA	LVS	114	14.151	2.356	3.550	1.00	0.00
ATOM	1691	CA	LVS	103	-0.056	6.514	3.900	1.00	0.00	ATOM	1785	N	SEK	108	5.411	7.766	5.883	1.00	0.00	ATOM	1879	HE1	LVS	113	12.128	1.872	3.167	1.00	0.00
ATOM	1692	CA	LVS	103	-1.446	8.146	3.857	1.00	0.00	ATOM	1786	N	SEK	108	4.619	7.305	6.227	1.00	0.00	ATOM	1880	HE2	LVS	113	12.018	1.381	4.123	1.00	0.00
ATOM	1693	CA	LVS	103	-1.446	8.146	3.857	1.00	0.00	ATOM	1787	CA	SEK	108	5.321	8.130	5.133	1.00	0.00	ATOM	1881	HE2	LVS	113	12.018	1.381	4.123	1.00	0.00
ATOM	1694	HE2	LVS	103	-2.220	7.612	3.122	1.00	0.00	ATOM	1788	CA	SEK	108	5.321	8.130	5.133	1.00	0.00	ATOM	1882	CA	LVS	113	13.513	1.715	1.873	1.00	0.00
ATOM	1695	CA	LVS	103	-0.670	8.991	2.860	1.00	0.00	ATOM	1789	CA	SEK	108	5.730	8.643	8.157	1.00	0.00	ATOM	1883	C	LVS	113	13.917	2.823	0.936	1.00	0.00
ATOM	1696	HE2	LVS	103	-1.083	8.832	1.874	1.00	0.00	ATOM	1790	HE1	SEK	108	4.648	8.597	8.942	1.00	0.00	ATOM	1884	N	LVS	114	13.917	2.823	0.936	1.00	0.00
ATOM	1697	HE2	LVS	103	0.364	8.680	2.871	1.00	0.00	ATOM	1791	HE2	SEK	108	4.940	7.938	8.164	1.00	0.00	ATOM	1885	N	LVS	114	13.917	2.823	0.936	1.00	0.00
ATOM	1698	CA	LVS	103	-0.734	10.472	3.178	1.00	0.00	ATOM	1792	CA	SEK	108	5.181	9.949	8.121	1.00	0.00	ATOM	1886	CA	LVS	114	13.917	2.823	0.936	1.00	0.00
ATOM	1699	HE2	LVS	103	-1.955	10.956	3.442	1.00	0.00	ATOM	1793	CA	SEK	108	4.501	9.993	7.445	1.00	0.00	ATOM	1887	HE1	LVS	114	14.518	5.484	0.228	1.00	0.00
ATOM	1700	CA	LVS	103	-1.955	10.956	3.442	1.00	0.00	ATOM	1794	C	SEK	108	7.149	7.561	7.212	1.00	0.00	ATOM	1888	C	LVS	114	14.518	5.484	0.228	1.00	0.00
ATOM	1701	C	LVS	103	0.439	7.858	2.474	1.00	0.00	ATOM	1795	C	SEK	108	7.149	7.561	7.212	1.00	0.00	ATOM	1889	C	LVS	114	14.518	5.484	0.228	1.00	0.00
ATOM	1702	C	LVS	103	1.644	7.658	5.334	1.00	0.00	ATOM	1796	N	LVS	109	7.049	5.966	6.594	1.00	0.00	ATOM	1890	C	LVS	114	13.227	4.056	-1.417	1.00	0.00
ATOM	1703	N	LVS	104	-0.052	8.675	6.405	1.00	0.00	ATOM	1797	N	LVS	109	6.100	5.811	6.828	1.00	0.00	ATOM	1891	N	LVS	114	13.227	4.056	-1.417	1.00	0.00
ATOM	1704	CA	LVS	104	-1.024	8.786	6.480	1.00	0.00	ATOM	1798	CA	LVS	109	7.953	4.860	7.124	1.00	0.00	ATOM	1892	CA	LVS	115	11.006	4.534	-0.031	1.00	0.00
ATOM	1705	CA	LVS	104	0.821	9.417	7.106	1.00	0.00	ATOM	1799	CA	LVS	109	8.630	5.055	7.951	1.00	0.00	ATOM	1893	CA	LVS	115	11.006	4.534	-0.031	1.00	0.00
ATOM	1706	CA	LVS	104	-0.289	10.172	6.255	1.00	0.00	ATOM	1800	CA	LVS	109	7.149	3.591	7.461	1.00	0.00	ATOM	1894	CA	LVS	115	11.132	4.180	-2.510	1.00	0.00
ATOM	1707	CA	LVS	104	-0.289	10.172	6.255	1.00	0.00	ATOM	1801	CA	LVS	109	7.149	3.591	7.461	1.00	0.00	ATOM	1895	CA	LVS	115	11.132	4.180	-2.510	1.00	0.00
ATOM	1708	HE1	LVS	104	-0.949	9.589	4.492	1.00	0.00	ATOM	1802	HE2	LVS	109	7.741	2.737	7.126	1.00	0.00	ATOM	1896	HE1	LVS	115	9.166	3.166	-0.144	1.00	0.00
ATOM	1709	HE1	LVS	104	0.530	10.034	9.335	1.00	0.00	ATOM	1803	CA	LVS	109	6.734	4.400	8.856	1.00	0.00	ATOM	1897	HE2	LVS	115	8.890	3.167	-1.754	1.00	0.00
ATOM	1710	CA	LVS	104	-0.278	11.559	8.125	1.00	0.00	ATOM	1804	HE1	LVS	109	5.820	2.866	8.902	1.00	0.00	ATOM	1898	HE2	LVS	115	10.497	2.058	-0.907	1.00	0.00
ATOM	1711	HE1	LVS	104	0.635	12.118	8.261	1.00	0.00	ATOM	1805	HE2	LVS	109	6.566	4.421	9.276	1.00	0.00	ATOM	1899	CA	LVS	115	9.789	2.266	-0.391	1.00	0.00
ATOM	1712	HE2	LVS	104	-1.028	11.950	8.820	1.00	0.00	ATOM	1806	CA	LVS	109	7.802	2.732	9.267	1.00	0.00	ATOM	1900	CA	LVS	115	9.789	2.266	-0.391	1.00	0.00
ATOM	1713	CA	LVS	104	-0.9																								

REPLACEMENT SHEET

ATCM	1947	HE2 LVS	118	4.432	9.001	-5.028	1.00	0.00
ATCM	1948	CG LVS	118	3.021	10.506	-5.527	1.00	0.00
ATCM	1949	HG1 LVS	118	3.175	10.140	-6.531	1.00	0.00
ATCM	1950	HG2 LVS	118	3.077	11.585	-5.522	1.00	0.00
ATCM	1951	HG3 LVS	118	1.422	9.100	-3.443	1.00	0.00
ATCM	1952	HDI LVS	118	1.422	9.100	-3.443	1.00	0.00
ATCM	1953	HDI LVS	118	1.422	9.100	-3.443	1.00	0.00
ATCM	1954	CE LVS	118	0.569	11.052	-5.518	1.00	0.00
ATCM	1955	HE1 LVS	118	-0.025	11.347	-4.666	1.00	0.00
ATCM	1956	HE2 LVS	118	1.048	11.922	-5.942	1.00	0.00
ATCM	1957	NZ LVS	118	-0.324	10.443	-6.543	1.00	0.00
ATCM	1958	HE3 LVS	118	-0.324	10.443	-6.543	1.00	0.00
ATCM	1959	HZ2 LVS	118	0.214	10.214	-7.468	1.00	0.00
ATCM	1960	HZ3 LVS	118	-1.097	11.039	-6.778	1.00	0.00
ATCM	1961	C LVS	118	4.974	12.103	-3.687	1.00	0.00
ATCM	1962	OT1 LVS	118	4.769	13.177	-4.291	1.00	0.00
ATCM	1963	OT2 LVS	118	4.901	11.986	-2.445	1.00	0.00
END								

Table 6

Atomic Structure Coordinates of the P/CAF Bromodomain/Acetyl-Histamine Complex

REMARK										
PDBNAME=7/blocho/chr16/complex_xplor/structure/tb/complex_81-REMARK										
initial random number seed: 5.960359E+10REMARK										
overall bond, angle, improper, vdw, no, cdhbmak emr(41, 107, 9, 85656,										
72.1621, 0, 22.2303, 36.0151, 0.204542REMARK										
bond, angle, improper, no, cdhbmak emr-dw: 2.214961E-										
03.0.361077.53.7899, 1.40651E-02.0.249335REMARK										
no, cdhbmak violations: 2, 0REMARK										
1.912	-4.558	1.00	0.00	ACH ATOM	2	HA1 ACE	200	-14.803	0.08	
2.641	-1.705	1.00	0.00	ACH ATOM	3	HA2 ACE	200	-13.320	0.00	
1.912	-4.558	1.00	0.00	ACH ATOM	4	HA3 ACE	200	-14.481	0.00	
2.304	-5.428	1.00	0.00	ACH ATOM	5	CAE	200	-13.332	0.00	
1.972	-5.067	1.00	0.00	ACH ATOM	6	CAE	200	-13.385	0.00	
2.936	-3.376	1.00	0.00	ACH ATOM	7	N	HIM	201	-13.773	0.00
2.238	-3.012	1.00	0.00	ACH ATOM	8	N	HIM	201	-13.773	0.00
4.957	-2.683	1.00	0.00	ACH ATOM	9	CN	HIM	201	-13.773	0.00
3.827	-1.417	1.00	0.00	ACH ATOM	10	HA1 HIM	201	-14.831	0.00	
3.589	-0.620	1.00	0.00	ACH ATOM	11	HA2 HIM	201	-15.236	0.00	
3.025	-1.536	1.00	0.00	ACH ATOM	12	CB HIM	201	-15.268	0.00	
5.100	-1.055	1.00	0.00	ACH ATOM	13	CB HIM	201	-15.591	0.00	
5.940	-1.107	1.00	0.00	ACH ATOM	14	H2 HIM	201	-15.663	0.00	
5.329	-2.010	1.00	0.00	ACH ATOM	15	NH1 HIM	201	-17.132	0.00	
6.295	-1.057	1.00	0.00	ACH ATOM	16	NH2 HIM	201	-17.421	0.00	
6.906	-1.057	1.00	0.00	ACH ATOM	17	H2 HIM	201	-17.421	0.00	
4.756	-3.204	1.00	0.00	ACH ATOM	18	CD HIM	201	-16.680	0.00	
4.116	-3.779	1.00	0.00	ACH ATOM	19	HD HIM	201	-16.516	0.00	
6.135	-2.729	1.00	0.00	ACH ATOM	20	CE HIM	201	-15.220	0.00	
5.321	-3.592	1.00	0.00	ACH ATOM	21	HE HIM	201	-15.220	0.00	
16.168	-0.871	1.00	0.00	ACH ATOM	22	CE HIM	201	-15.220	0.00	
15.945	-0.355	1.00	0.00	ACH ATOM	23	CA GUY	1	28.103	0.00	
17.114	1.011	1.00	0.00	ACH ATOM	24	HA1 GUY	1	28.724	0.00	
17.834	1.144	1.00	0.00	ACH ATOM	25	CA GUY	1	28.713	0.00	
17.963	-1.018	1.00	0.00	ACH ATOM	26	C GUY	1	26.700	0.00	
17.584	-2.207	1.00	0.00	ACH ATOM	27	O GUY	1	26.911	0.00	
18.667	-0.970	1.00	0.00	ACH ATOM	28	N GUY	1	27.307	0.00	
16.682	2.037	1.00	0.00	ACH ATOM	29	PT GUY	1	26.999	0.00	
16.110	1.866	1.00	0.00	ACH ATOM	30	N SER	2	27.322	0.00	
17.043	3.411	1.00	0.00	ACH ATOM	31	CA SER	2	27.328	0.00	
17.395	3.975	1.00	0.00	ACH ATOM	32	HA SER	2	28.283	0.00	
18.588	4.762	1.00	0.00	ACH ATOM	33	HA SER	2	28.283	0.00	
17.414	4.374	1.00	0.00	ACH ATOM	34	HA SER	2	28.283	0.00	
18.897	2.969	1.00	0.00	ACH ATOM	35	HA SER	2	28.283	0.00	
19.435	2.650	1.00	0.00	ACH ATOM	36	HA SER	2	28.283	0.00	
15.801	4.250	1.00	0.00	ACH ATOM	37	HA SER	2	28.283	0.00	
15.803	5.415	1.00	0.00	ACH ATOM	38	HA SER	2	28.283	0.00	
16.803	5.627	1.00	0.00	ACH ATOM	39	HA SER	2	28.283	0.00	
14.824	6.342	1.00	0.00	ACH ATOM	40	HA SER	2	28.283	0.00	
13.972	5.775	1.00	0.00	ACH ATOM	41	HA SER	2	28.283	0.00	
15.176	7.399	1.00	0.00	ACH ATOM	42	HA SER	2	28.283	0.00	
14.438	8.178	1.00	0.00	ACH ATOM	43	HA SER	2	28.283	0.00	
14.438	8.178	1.00	0.00	ACH ATOM	44	HA SER	2	28.283	0.00	
16.718	9.062	1.00	0.00	ACH ATOM	45	HA SER	2	28.283	0.00	
16.718	9.062	1.00	0.00	ACH ATOM	46	HA SER	2	28.283	0.00	
17.975	9.465	1.00	0.00	ACH ATOM	47	HA SER	2	28.283	0.00	
17.975	9.465	1.00	0.00	ACH ATOM	48	HA SER	2	28.283	0.00	
18.971	10.150	1.00	0.00	ACH ATOM	49	HA SER	2	28.283	0.00	
18.615	8.668	1.00	0.00	ACH ATOM	50	HA SER	2	28.283	0.00	
18.615	8.668	1.00	0.00	ACH ATOM	51	HA SER	2	28.283	0.00	
14.663	7.020	1.00	0.00	ACH ATOM	52	HA SER	2	28.283	0.00	
14.998	8.078	1.00	0.00	ACH ATOM	53	HA SER	2	28.283	0.00	
13.551	6.005	1.00	0.00	ACH ATOM	54	HA SER	2	28.283	0.00	
13.119	6.351	1.00	0.00	ACH ATOM	55	HA SER	2	28.283	0.00	
12.767	7.958	1.00	0.00	ACH ATOM	56	HA SER	2	28.283	0.00	
14.294	6.994	1.00	0.00	ACH ATOM	57	HA SER	2	28.283	0.00	
14.294	6.994	1.00	0.00	ACH ATOM	58	HA SER	2	28.283	0.00	
13.962	5.981	1.00	0.00	ACH ATOM	59	HA SER	2	28.283	0.00	
14.176	8.727	1.00	0.00	ACH ATOM	60	HA SER	2	28.283	0.00	
12.911	7.535	1.00	0.00	ACH ATOM	61	HA SER	2	28.283	0.00	
14.909	7.006	1.00	0.00	ACH ATOM	62	HA SER	2	28.283	0.00	
16.531	6.842	1.00	0.00	ACH ATOM	63	HA SER	2	28.283	0.00	
15.643	7.000	1.00	0.00	ACH ATOM	64	HA SER	2	28.283	0.00	
16.638	5.851	1.00	0.00	ACH ATOM	65	HA SER	2	28.283	0.00	
11.977	6.126	1.00	0.00	ACH ATOM	66	HA SER	2	28.283	0.00	
11.906	4.914	1.00	0.00	ACH ATOM	67	HA SER	2	28.283	0.00	
11.084	6.792	1.00	0.00	ACH ATOM	68	HA SER	2	28.283	0.00	
11.134	7.758	1.00	0.00	ACH ATOM	69	HA SER	2	28.283	0.00	
9.939	6.425	1.00	0.00	ACH ATOM	70	HA SER	2	28.283	0.00	
9.560	7.123	1.00	0.00	ACH ATOM	71	HA SER	2	28.283	0.00	
8.114	6.801	1.00	0.00	ACH ATOM	72	HA SER	2	28.283	0.00	
9.210	8.421	1.00	0.00	ACH ATOM	73	HA SER	2	28.283	0.00	
8.428	8.588	1.00	0.00	ACH ATOM	74	HA SER	2	28.283	0.00	
10.380	4.934	1.00	0.00	ACH ATOM	75	HA SER	2	28.283	0.00	
10.380	4.934	1.00	0.00	ACH ATOM	76	HA SER	2	28.283	0.00	
9.465	4.110	1.00	0.00	ACH ATOM	77	HA SER	2	28.283	0.00	
8.565	4.110	1.00	0.00	ACH ATOM	78	HA SER	2	28.283	0.00	
9.753	2.911	1.00	0.00	ACH ATOM	79	HA SER	2	28.283	0.00	
10.720	2.950	1.00	0.00	ACH ATOM	80	HA SER	2	28.283	0.00	
9.801	1.572	1.00	0.00	ACH ATOM	81	HA SER	2	28.283	0.00	
10.856	1.572	1.00	0.00	ACH ATOM	82	HA SER	2	28.283	0.00	
10.856	1.572	1.00	0.00	ACH ATOM	83	HA SER	2	28.283	0.00	
10.856	1.572	1.00	0.00	ACH ATOM	84	HA SER	2	28.283	0.00	
10.856	1.572	1.00	0.00	ACH ATOM	85	HA SER	2	28.283	0.00	
10.856	1.572	1.00	0.00	ACH ATOM	86	HA SER	2	28.283	0.00	
10.856	1.572	1.00	0.00	ACH ATOM	87	HA SER	2	28.283	0.00	
10.856	1.572	1.00	0.00	ACH ATOM	88	HA SER	2	28.283	0.00	
10.856	1.572	1.00	0.00	ACH ATOM	89	HA SER	2	28.283	0.00	
10.856	1.572	1.00	0.00	ACH ATOM	90	HA SER	2	28.283	0.00	
10.856	1.572	1.00	0.00	ACH ATOM	91	HA SER	2	28.283	0.00	
10.856	1.572	1.00	0.00	ACH ATOM	92	HA SER	2	28.283	0.00	
10.856	1.572	1.00	0.00	ACH ATOM	93	HA SER	2	28.283	0.00	
10.856	1.572	1.00	0.00	ACH ATOM	94	HA SER	2	28.283	0.00	
10.856	1.572	1.00	0.00	ACH ATOM	95	HA SER	2	28.283	0.00	
10.856	1.572	1.00	0.00	ACH ATOM	96	HA SER	2	28.283	0.00	
10.856	1.572	1.00	0.00	ACH ATOM	97	HA SER	2	28.283	0.00	
10.856	1.572	1.00	0.00	ACH ATOM	98	HA SER	2	28.283	0.00	
10.856	1.572	1.00	0.00	ACH ATOM	99	HA SER	2	28.283	0.00	
10.856	1.572	1.00	0.00	ACH ATOM	100	HA SER	2	28.283	0.00	
10.856	1.572	1.00	0.00	ACH ATOM	101	HA SER	2	28.283	0.00	
10.856	1.572	1.00	0.00	ACH ATOM	102	HA SER	2	28.283	0.00	
10.856	1.572	1.00	0.00	ACH ATOM	103	HA SER	2	28.283	0.00	
10.856	1.572	1.00	0.00	ACH ATOM	104	HA SER	2	28.283	0.00	
10.856	1.572	1.00	0.00	ACH ATOM	105	HA SER	2	28.283	0.00	
10.856	1.572	1.00	0.00	ACH ATOM	106	HA SER	2	28.283	0.00	
10.856	1.572	1.00	0.00	ACH ATOM	107	HA SER	2	28.283	0.00	
10.856	1.572	1.00	0.00	ACH ATOM	108	HA SER	2	28.283	0.00	
10.856	1.572	1.00	0.00	ACH ATOM	109	HA SER	2	28.283	0.00	
10.856	1.572	1.00	0.00	ACH ATOM	110	HA SER	2	28.283	0.00	
10.856	1.572	1.00	0.00	ACH ATOM	111	HA SER	2	28.283	0.00	
10.856	1.572	1.00	0.00	ACH ATOM	112	HA SER	2	28.283	0.00	
10.856	1.572	1.00	0.00	ACH ATOM	113	HA SER	2	28.283	0.00	
10.856	1.572	1.00	0.00	ACH ATOM	114	HA SER	2	28.283	0.00	
10.856	1.572	1.00	0.00	ACH ATOM	115	HA SER	2	28.283	0.00	
10.856	1.572	1.00	0.00	ACH ATOM	116	HA SER	2	28.283	0.00	
10.856	1.572	1.00	0.00	ACH ATOM	117	HA SER	2	28.283	0.00	
10.856	1.572	1.00	0.00	ACH ATOM	118	HA SER	2	28.283	0.00	
10.856	1.572	1.00	0.00	ACH ATOM	119	HA SER	2	28.283		

REPLACEMENT SHEET

7.249	-6.857	1.00	0.00	BRD ATOM	253	HE2	TYR	15	9.275	8.433	2.377	1.00	0.00	BRD ATOM	347	HG22	TLE	21	3.319	12.845	-9.465	1.00	0.00	BRD ATOM	443	HE1	LVS	26	-1.215
7.326	-7.111	1.00	0.00	BRD ATOM	254	CE	TYR	15	10.908	6.933	2.603	1.00	0.00	BRD ATOM	348	HG23	TLE	21	3.073	13.285	-9.484	1.00	0.00	BRD ATOM	443	HE2	LVS	26	-0.010
7.352	-7.706	1.00	0.00	BRD ATOM	255	HH	TYR	15	10.908	7.915	3.053	1.00	0.00	BRD ATOM	349	COL	TLE	21	3.222	13.461	-8.710	1.00	0.00	BRD ATOM	444	CE	LVS	26	-0.358
8.358	-8.883	1.00	0.00	BRD ATOM	256	HH	TYR	15	10.972	6.140	-0.427	1.00	0.00	BRD ATOM	350	HDI1	TLE	21	3.805	12.749	-10.199	1.00	0.00	BRD ATOM	444	CE	LVS	26	-0.673
8.367	-8.997	1.00	0.00	BRD ATOM	257	C	TYR	15	10.959	6.000	-0.338	1.00	0.00	BRD ATOM	351	HDI2	TLE	21	2.284	11.073	-5.468	1.00	0.00	BRD ATOM	445	O	LVS	27	-4.762
7.230	-8.977	1.00	0.00	BRD ATOM	257	C	TYR	15	10.959	6.000	-0.338	1.00	0.00	BRD ATOM	352	HDI3	TLE	21	3.075	10.991	-6.056	1.00	0.00	BRD ATOM	446	N	SER	27	-4.315
7.320	-8.977	1.00	0.00	BRD ATOM	258	N	SER	16	11.678	9.361	-1.044	1.00	0.00	BRD ATOM	353	C	TLE	21	1.989	12.116	-4.732	1.00	0.00	BRD ATOM	447	HA	SER	27	-6.104
8.026	-1.146	1.00	0.00	BRD ATOM	259	HA	SER	16	12.485	9.361	-1.044	1.00	0.00	BRD ATOM	354	N	TLE	21	2.433	13.257	-4.516	1.00	0.00	BRD ATOM	448	HA	SER	27	-6.104
7.666	-1.605	1.00	0.00	BRD ATOM	260	HA	SER	16	11.249	8.608	-2.034	1.00	0.00	BRD ATOM	355	N	LVS	22	3.325	13.100	-5.091	1.00	0.00	BRD ATOM	449	HA	SER	27	-4.532
9.384	-1.602	1.00	0.00	BRD ATOM	261	HA	SER	16	11.148	8.608	-2.034	1.00	0.00	BRD ATOM	356	N	LVS	22	1.668	14.553	-4.976	1.00	0.00	BRD ATOM	450	HA	SER	27	-5.645
9.346	-2.518	1.00	0.00	BRD ATOM	262	HA	SER	16	11.148	8.608	-2.034	1.00	0.00	BRD ATOM	357	CA	LVS	22	0.757	15.781	-4.404	1.00	0.00	BRD ATOM	451	HA	SER	27	-5.207
9.040	-2.518	1.00	0.00	BRD ATOM	263	CA	SER	16	13.505	8.317	-3.229	1.00	0.00	BRD ATOM	358	HA	LVS	22	2.451	15.781	-4.404	1.00	0.00	BRD ATOM	452	HA	SER	27	-5.207
9.040	-2.518	1.00	0.00	BRD ATOM	264	HA	SER	16	13.505	8.317	-3.229	1.00	0.00	BRD ATOM	359	HA	LVS	22	2.451	15.781	-4.404	1.00	0.00	BRD ATOM	453	HA	SER	27	-5.207
10.185	-1.288	1.00	0.00	BRD ATOM	265	HA	SER	16	12.601	7.807	-2.929	1.00	0.00	BRD ATOM	360	HA	LVS	22	2.132	14.462	-3.551	1.00	0.00	BRD ATOM	454	HA	SER	27	-3.746
11.317	-0.405	1.00	0.00	BRD ATOM	266	HA	SER	16	12.601	7.807	-2.929	1.00	0.00	BRD ATOM	361	HA	LVS	22	1.772	14.462	-3.551	1.00	0.00	BRD ATOM	455	HA	SER	27	-3.746
11.992	-0.405	1.00	0.00	BRD ATOM	267	HA	SER	16	9.909	7.986	-4.462	1.00	0.00	BRD ATOM	362	HA	LVS	22	1.772	14.462	-3.551	1.00	0.00	BRD ATOM	456	HA	SER	27	-3.746
9.370	-0.879	1.00	0.00	BRD ATOM	268	C	SER	16	9.028	7.082	-5.486	1.00	0.00	BRD ATOM	363	HA	LVS	22	1.378	14.462	-3.551	1.00	0.00	BRD ATOM	457	HA	SER	27	-3.746
10.182	-1.161	1.00	0.00	BRD ATOM	269	C	SER	16	9.028	7.082	-5.486	1.00	0.00	BRD ATOM	364	COL	LVS	22	2.665	12.426	-2.339	1.00	0.00	BRD ATOM	458	HA	SER	27	-5.215
8.420	0.038	1.00	0.00	BRD ATOM	270	N	TYR	17	10.493	6.362	-6.437	1.00	0.00	BRD ATOM	365	HDI1	LVS	22	2.220	11.418	-2.785	1.00	0.00	BRD ATOM	459	HA	SER	27	-5.863
8.420	0.038	1.00	0.00	BRD ATOM	271	HA	TYR	17	10.493	6.362	-6.437	1.00	0.00	BRD ATOM	366	HDI2	LVS	22	3.601	12.137	-0.946	1.00	0.00	BRD ATOM	460	HA	SER	27	-5.597
8.420	0.038	1.00	0.00	BRD ATOM	272	HA	TYR	17	10.493	6.362	-6.437	1.00	0.00	BRD ATOM	367	HDI3	LVS	22	2.845	11.312	-0.497	1.00	0.00	BRD ATOM	461	HA	SER	27	-5.125
8.268	0.785	1.00	0.00	BRD ATOM	273	CB	TYR	17	8.656	5.361	-6.072	1.00	0.00	BRD ATOM	368	HDI1	LVS	22	0.137	13.182	-0.497	1.00	0.00	BRD ATOM	462	HA	SER	27	-4.283
7.146	1.820	1.00	0.00	BRD ATOM	274	CB	TYR	17	9.069	6.222	-5.178	1.00	0.00	BRD ATOM	369	HDI2	LVS	22	-0.259	10.146	-0.742	1.00	0.00	BRD ATOM	463	HA	SER	27	-5.540
6.212	-1.197	1.00	0.00	BRD ATOM	275	HB	TYR	17	10.426	6.401	-5.199	1.00	0.00	BRD ATOM	370	HDI2	LVS	22	0.790	10.086	1.205	1.00	0.00	BRD ATOM	464	HA	SER	27	-5.540
5.282	-2.591	1.00	0.00	BRD ATOM	276	COI	TYR	17	10.426	6.401	-5.199	1.00	0.00	BRD ATOM	371	HDI3	LVS	22	1.243	11.931	2.081	1.00	0.00	BRD ATOM	465	HA	SER	27	-5.540
5.424	-3.243	1.00	0.00	BRD ATOM	277	COI	TYR	17	10.426	6.401	-5.199	1.00	0.00	BRD ATOM	372	C	LVS	22	0.086	12.883	1.866	1.00	0.00	BRD ATOM	466	HA	SER	27	-5.540
4.868	-3.198	1.00	0.00	BRD ATOM	278	COI	TYR	17	7.346	9.636	-3.915	1.00	0.00	BRD ATOM	373	C	LVS	22	2.201	9.754	1.966	1.00	0.00	BRD ATOM	467	HA	SER	27	-5.540
4.356	-1.720	1.00	0.00	BRD ATOM	279	COI	TYR	17	6.714	9.636	-3.915	1.00	0.00	BRD ATOM	374	C	LVS	22	2.201	9.754	1.966	1.00	0.00	BRD ATOM	468	HA	SER	27	-5.540
4.356	-1.720	1.00	0.00	BRD ATOM	280	COI	TYR	17	6.714	9.636	-3.915	1.00	0.00	BRD ATOM	375	C	LVS	22	2.201	9.754	1.966	1.00	0.00	BRD ATOM	469	HA	SER	27	-5.540
2.840	-2.622	1.00	0.00	BRD ATOM	281	COI	TYR	17	7.546	10.354	-3.725	1.00	0.00	BRD ATOM	376	CA	LVS	23	1.941	11.466	3.262	1.00	0.00	BRD ATOM	470	HA	SER	27	-6.314
2.893	-3.455	1.00	0.00	BRD ATOM	282	C	TYR	17	7.356	10.354	-3.725	1.00	0.00	BRD ATOM	377	CA	LVS	23	1.941	11.466	3.262	1.00	0.00	BRD ATOM	471	HA	SER	27	-6.314
7.971	-0.161	1.00	0.00	BRD ATOM	283	C	TYR	17	6.420	11.670	-5.754	1.00	0.00	BRD ATOM	378	CA	LVS	23	3.118	12.044	4.141	1.00	0.00	BRD ATOM	472	HA	SER	27	-6.314
8.757	-0.277	1.00	0.00	BRD ATOM	284	N	LVS	18	7.438	11.670	-5.754	1.00	0.00	BRD ATOM	379	HA	LVS	23	3.720	9.536	3.925	1.00	0.00	BRD ATOM	473	C	HIS	28	-7.364
6.932	-0.843	1.00	0.00	BRD ATOM	285	N	LVS	18	8.212	12.793	-6.453	1.00	0.00	BRD ATOM	380	HA	LVS	23	2.731	12.017	-0.850	1.00	0.00	BRD ATOM	474	N	HIS	28	-7.979
6.932	-0.843	1.00	0.00	BRD ATOM	286	HA	LVS	18	5.356	12.793	-6.453	1.00	0.00	BRD ATOM	381	HA	LVS	23	4.012	11.017	-0.850	1.00	0.00	BRD ATOM	475	N	HIS	28	-7.979
6.414	-1.776	1.00	0.00	BRD ATOM	287	HA	LVS	18	6.871	11.969	-6.035	1.00	0.00	BRD ATOM	382	HA	LVS	23	4.886	11.386	-0.850	1.00	0.00	BRD ATOM	476	HA	LVS	28	-6.930
5.122	-1.197	1.00	0.00	BRD ATOM	288	CB	LVS	18	7.669	11.969	-6.035	1.00	0.00	BRD ATOM	383	HDI2	LVS	23	4.886	12.608	-0.687	1.00	0.00	BRD ATOM	477	HA	LVS	28	-6.930
5.282	-2.591	1.00	0.00	BRD ATOM	289	HB	LVS	18	6.439	13.535	-5.393	1.00	0.00	BRD ATOM	384	CB	LVS	23	3.316	12.608	-0.687	1.00	0.00	BRD ATOM	478	HA	LVS	28	-6.930
5.424	-3.243	1.00	0.00	BRD ATOM	290	CG	LVS	18	7.377	13.697	-6.815	1.00	0.00	BRD ATOM	385	CG	LVS	23	2.634	12.608	-0.687	1.00	0.00	BRD ATOM	479	HA	LVS	28	-6.930
4.868	-3.198	1.00	0.00	BRD ATOM	291	CG	LVS	18	7.377	13.697	-6.815	1.00	0.00	BRD ATOM	386	CG	LVS	23	3.494	13.800	-0.687	1.00	0.00	BRD ATOM	480	HA	LVS	28	-6.930
4.356	-1.720	1.00	0.00	BRD ATOM	292	HA	LVS	18	8.301	14.692	-8.034	1.00	0.00	BRD ATOM	387	HA	LVS	23	4.058	13.800	-0.687	1.00	0.00	BRD ATOM	481	HA	LVS	28	-6.930
4.356	-1.720	1.00	0.00	BRD ATOM	293	HA	LVS	18	8.301	14.692	-8.034	1.00	0.00	BRD ATOM	388	HA	LVS	23	3.052	13.901	1.092	1.00	0.00	BRD ATOM	482	HA	LVS	28	-6.930
2.840	-2.622	1.00	0.00	BRD ATOM	294	HDI1	LVS	18	8.667	13.464	-4.157	1.00	0.00	BRD ATOM	389	HDI2	LVS	23	0.207	15.165	0.093	1.00	0.00	BRD ATOM	483	HA	LVS	28	-6.930
2.893	-3.455	1.00	0.00	BRD ATOM	295	HDI2	LVS	18	7.523	12.692	-4.925	1.00	0.00	BRD ATOM	390	C	LVS	23	0.530	16.105	0.093	1.00	0.00	BRD ATOM	484	HA	LVS	28	-6.930
1.941	-2.039	1.00	0.00	BRD ATOM	297	COI	LVS	18	6.439	12.592	-2.833	1.00	0.00	BRD ATOM	391	N	LVS	24	1.252	15.165	0.093	1.00	0.00	BRD ATOM	485	HA	LVS	28	-6.930
3.727	-0.658	1.00	0.00	BRD ATOM	298	HDI1	LVS	18	6.439	12.592	-2.833	1.00	0.00	BRD ATOM	392	N	LVS	24	1.252	15.165	0.093	1.00	0.00	BRD ATOM	486	HA	LVS	28	-6.930
4.180	-0.897	1.00	0.00	BRD ATOM	299	HDI2	LVS	18	6.252	12.167	-2.269	1.00	0.00	BRD ATOM	393	CA	LVS	24	-0.654	16.290	-0.996	1.00	0.00	BRD ATOM	487	HA	LVS	28	-6.930
4.180	-0.897	1.00	0.00	BRD ATOM	300	HDI2	LVS	18	6.252	12.167	-2.269	1.00	0.00	BRD AT															

9.152	-0.365	1.00	0.00	BD ATOM	817	HB	11E	50	-6.001	0.533	-6.566	1.00	0.00	BD ATOM	911	CA	LEU	56	-3.106	5.934	-13.617	1.00	0.00	BD ATOM	1005	C	GLU	61	3.565	-
9.428	-0.635	1.00	0.00	BD ATOM	818	CO1	11E	50	-7.818	2.010	-6.111	1.00	0.00	BD ATOM	912	CA	LEU	56	-1.300	-1.400	-11.400	1.00	0.00	BD ATOM	1006	N	ARG	62	4.456	-
9.035	0.436	1.00	0.00	BD ATOM	819	HO1	11E	50	-8.256	1.790	-5.810	1.00	0.00	BD ATOM	913	HB	LEU	56	-3.804	3.601	-11.782	1.00	0.00	BD ATOM	1007	N	ARG	62	4.456	-
10.011	0.288	1.00	0.00	BD ATOM	820	HO2	11E	50	-8.158	3.224	-5.292	1.00	0.00	BD ATOM	914	HB	LEU	56	-3.692	2.137	-11.244	1.00	0.00	BD ATOM	1008	N	ARG	62	2.971	-
8.717	-0.344	1.00	0.00	BD ATOM	821	CO2	11E	50	-5.644	4.115	-5.092	1.00	0.00	BD ATOM	915	HB	LEU	56	-4.952	1.838	-10.697	1.00	0.00	BD ATOM	1009	CA	ARG	62	4.999	-
8.434	0.644	1.00	0.00	BD ATOM	822	HO2	11E	50	-6.289	3.082	-5.089	1.00	0.00	BD ATOM	916	CA	LEU	56	-3.088	1.428	-11.155	1.00	0.00	BD ATOM	1010	CA	ARG	62	5.352	-
7.912	-0.344	1.00	0.00	BD ATOM	823	HO2	11E	50	-6.289	3.451	-3.292	1.00	0.00	BD ATOM	917	HO	LEU	56	-3.356	0.975	-10.468	1.00	0.00	BD ATOM	1011	HB	ARG	62	4.807	-
7.250	1.139	1.00	0.00	BD ATOM	824	HO2	11E	50	-8.324	4.770	-3.344	1.00	0.00	BD ATOM	918	HO1	LEU	56	-4.497	0.187	-9.668	1.00	0.00	BD ATOM	1012	HB	ARG	62	4.051	-
7.981	1.169	1.00	0.00	BD ATOM	825	CO1	11E	50	-8.330	5.539	-3.606	1.00	0.00	BD ATOM	919	HO1	LEU	56	-4.497	1.018	-11.404	1.00	0.00	BD ATOM	1013	HO1	ARG	62	4.832	-
8.499	1.757	1.00	0.00	BD ATOM	826	HO1	11E	50	-7.674	5.040	-3.719	1.00	0.00	BD ATOM	920	HO1	LEU	56	-3.562	0.928	-12.424	1.00	0.00	BD ATOM	1014	HO1	ARG	62	4.494	-
9.393	2.489	1.00	0.00	BD ATOM	827	HO2	11E	50	-9.326	4.668	-2.270	1.00	0.00	BD ATOM	921	CO2	LEU	56	-1.350	1.928	-11.495	1.00	0.00	BD ATOM	1015	HO2	ARG	62	4.832	-
7.704	2.094	1.00	0.00	BD ATOM	828	HO1	11E	50	-4.490	3.419	-4.153	1.00	0.00	BD ATOM	922	CO2	LEU	56	-1.144	1.278	-11.225	1.00	0.00	BD ATOM	1016	HO2	ARG	62	2.484	-
8.116	1.631	1.00	0.00	BD ATOM	829	C	11E	50	-4.504	3.188	-5.209	1.00	0.00	BD ATOM	923	HO2	LEU	56	-1.105	0.738	-10.355	1.00	0.00	BD ATOM	1017	HO2	ARG	62	2.652	-
11.444	-0.327	1.00	0.00	BD ATOM	830	N	ARG	51	-3.443	2.562	-3.671	1.00	0.00	BD ATOM	924	HO2	LEU	56	-1.105	0.851	-12.382	1.00	0.00	BD ATOM	1018	HO2	ARG	62	2.052	-
10.597	1.428	1.00	0.00	BD ATOM	831	CA	ARG	51	-2.082	2.400	-7.587	1.00	0.00	BD ATOM	925	N	LEU	56	-4.475	0.028	-12.346	1.00	0.00	BD ATOM	1019	CA	ARG	62	1.903	-
10.086	2.260	1.00	0.00	BD ATOM	832	CA	ARG	51	-2.082	2.760	-8.150	1.00	0.00	BD ATOM	926	N	LEU	56	-4.475	0.028	-12.346	1.00	0.00	BD ATOM	1020	CA	ARG	62	1.903	-
11.061	0.969	1.00	0.00	BD ATOM	833	HA	ARG	51	-0.968	2.304	-8.207	1.00	0.00	BD ATOM	927	HA	LEU	57	-4.534	1.585	-13.485	1.00	0.00	BD ATOM	1021	HA	ARG	62	1.183	-
12.139	1.024	1.00	0.00	BD ATOM	834	CA	ARG	51	-0.968	2.202	-7.696	1.00	0.00	BD ATOM	928	HA	LEU	57	-4.534	1.125	-14.498	1.00	0.00	BD ATOM	1022	CA	ARG	62	1.072	-
12.451	1.970	1.00	0.00	BD ATOM	835	HB	ARG	51	-1.452	2.639	-9.415	1.00	0.00	BD ATOM	929	HB	LEU	57	-4.534	0.828	-15.324	1.00	0.00	BD ATOM	1023	HB	ARG	62	2.473	-
10.481	1.341	1.00	0.00	BD ATOM	836	HB	ARG	51	-0.498	2.172	-10.108	1.00	0.00	BD ATOM	930	HB	LEU	57	-4.534	0.738	-10.355	1.00	0.00	BD ATOM	1024	HB	ARG	62	2.473	-
9.689	1.341	1.00	0.00	BD ATOM	837	CO2	ARG	51	-0.399	2.172	-10.108	1.00	0.00	BD ATOM	931	HA	LEU	57	-4.534	0.738	-10.355	1.00	0.00	BD ATOM	1025	HB2	ARG	62	2.473	-
11.516	2.301	1.00	0.00	BD ATOM	838	HO2	ARG	51	-0.399	2.172	-10.108	1.00	0.00	BD ATOM	932	HA	LEU	57	-4.534	0.738	-10.355	1.00	0.00	BD ATOM	1026	HB2	ARG	62	2.473	-
12.185	3.022	1.00	0.00	BD ATOM	839	HO2	ARG	51	-0.399	2.172	-10.108	1.00	0.00	BD ATOM	933	HA	LEU	57	-4.534	0.738	-10.355	1.00	0.00	BD ATOM	1027	HB2	ARG	62	2.473	-
11.005	2.753	1.00	0.00	BD ATOM	840	HO2	ARG	51	-0.399	2.172	-10.108	1.00	0.00	BD ATOM	934	HA	LEU	57	-4.534	0.738	-10.355	1.00	0.00	BD ATOM	1028	HB2	ARG	62	2.473	-
12.329	1.120	1.00	0.00	BD ATOM	841	CD	ARG	51	-0.552	2.328	-11.608	1.00	0.00	BD ATOM	935	CO	LEU	57	-5.133	2.606	-9.158	1.00	0.00	BD ATOM	1029	C	ARG	62	6.040	-
11.713	0.230	1.00	0.00	BD ATOM	842	HO2	ARG	51	-0.900	1.845	-12.118	1.00	0.00	BD ATOM	936	HO2	LEU	57	-6.248	2.328	-9.158	1.00	0.00	BD ATOM	1030	C	ARG	62	6.040	-
12.789	1.318	1.00	0.00	BD ATOM	843	HO2	ARG	51	-1.523	1.856	-11.889	1.00	0.00	BD ATOM	937	HO2	LEU	57	-6.248	2.328	-9.158	1.00	0.00	BD ATOM	1031	N	LEU	63	4.875	-
12.789	1.318	1.00	0.00	BD ATOM	844	HO2	ARG	51	-1.523	1.856	-11.889	1.00	0.00	BD ATOM	938	HO2	LEU	57	-6.248	2.328	-9.158	1.00	0.00	BD ATOM	1032	N	LEU	63	4.875	-
13.279	0.344	1.00	0.00	BD ATOM	845	CO2	ARG	51	2.683	4.068	-13.327	1.00	0.00	BD ATOM	939	HO2	LEU	57	-6.248	2.328	-9.158	1.00	0.00	BD ATOM	1033	CA	LEU	63	6.631	-
13.678	0.602	1.00	0.00	BD ATOM	846	CO2	ARG	51	2.683	4.068	-13.327	1.00	0.00	BD ATOM	940	CO2	LEU	57	-6.248	2.328	-9.158	1.00	0.00	BD ATOM	1034	CA	LEU	63	6.631	-
14.066	-0.132	1.00	0.00	BD ATOM	847	HO1	ARG	51	4.491	4.068	-13.327	1.00	0.00	BD ATOM	941	CO2	LEU	57	-6.248	2.328	-9.158	1.00	0.00	BD ATOM	1035	CA	LEU	63	6.631	-
13.610	1.532	1.00	0.00	BD ATOM	848	HO1	ARG	51	4.284	3.073	-13.697	1.00	0.00	BD ATOM	942	HO2	LEU	57	-6.778	4.277	-7.302	1.00	0.00	BD ATOM	1036	HB	LEU	63	5.009	-
13.610	1.532	1.00	0.00	BD ATOM	849	HO2	ARG	51	2.206	4.515	-12.715	1.00	0.00	BD ATOM	943	HO2	LEU	57	-6.778	2.962	-6.442	1.00	0.00	BD ATOM	1037	HB2	LEU	63	6.736	-
13.610	1.532	1.00	0.00	BD ATOM	850	HO2	ARG	51	2.206	4.515	-12.715	1.00	0.00	BD ATOM	944	HO2	LEU	57	-6.778	2.962	-6.442	1.00	0.00	BD ATOM	1038	CG	LEU	63	6.736	-
13.610	1.532	1.00	0.00	BD ATOM	851	HO2	ARG	51	2.206	4.515	-12.715	1.00	0.00	BD ATOM	945	HO2	LEU	57	-6.778	2.962	-6.442	1.00	0.00	BD ATOM	1039	CG	LEU	63	6.736	-
13.610	1.532	1.00	0.00	BD ATOM	852	HO2	ARG	51	2.206	4.515	-12.715	1.00	0.00	BD ATOM	946	HO2	LEU	57	-6.778	2.962	-6.442	1.00	0.00	BD ATOM	1040	CG	LEU	63	6.736	-
13.610	1.532	1.00	0.00	BD ATOM	853	HO2	ARG	51	2.206	4.515	-12.715	1.00	0.00	BD ATOM	947	HO2	LEU	57	-6.778	2.962	-6.442	1.00	0.00	BD ATOM	1041	HO1	LEU	63	8.237	-
13.610	1.532	1.00	0.00	BD ATOM	854	HO2	ARG	51	2.206	4.515	-12.715	1.00	0.00	BD ATOM	948	HO2	LEU	57	-6.778	2.962	-6.442	1.00	0.00	BD ATOM	1042	HO1	LEU	63	8.237	-
13.610	1.532	1.00	0.00	BD ATOM	855	HO2	ARG	51	2.206	4.515	-12.715	1.00	0.00	BD ATOM	949	HO2	LEU	57	-6.778	2.962	-6.442	1.00	0.00	BD ATOM	1043	HO1	LEU	63	8.237	-
13.610	1.532	1.00	0.00	BD ATOM	856	HO2	ARG	51	2.206	4.515	-12.715	1.00	0.00	BD ATOM	950	HO2	LEU	57	-6.778	2.962	-6.442	1.00	0.00	BD ATOM	1044	HO1	LEU	63	8.237	-
13.610	1.532	1.00	0.00	BD ATOM	857	HO2	ARG	51	2.206	4.515	-12.715	1.00	0.00	BD ATOM	951	HO2	LEU	57	-6.778	2.962	-6.442	1.00	0.00	BD ATOM	1045	HO1	LEU	63	8.237	-
13.610	1.532	1.00	0.00	BD ATOM	858	HO2	ARG	51	2.206	4.515	-12.715	1.00	0.00	BD ATOM	952	HO2	LEU	57	-6.778	2.962	-6.442	1.00	0.00	BD ATOM	1046	HO1	LEU	63	8.237	-
13.610	1.532	1.00	0.00	BD ATOM	859	HO2	ARG	51	2.206	4.515	-12.715	1.00	0.00	BD ATOM	953	HO2	LEU	57	-6.778	2.962	-6.442	1.00	0.00	BD ATOM	1047	HO1	LEU	63	8.237	-
13.610	1.532	1.00	0.00	BD ATOM	860	HO2	ARG	51	2.206	4.515	-12.715	1.00	0.00	BD ATOM	954	HO2	LEU	57	-6.778	2.962	-6.442	1.00	0.00	BD ATOM	1048	HO1	LEU	63	8.237	-
13.610	1.532	1.00	0.00	BD ATOM	861	HO2	ARG	51	2.206	4.515	-12.715	1.00	0.00	BD ATOM	955	HO2	LEU	57	-6.778	2.962	-6.442	1.00	0.00	BD ATOM	1049	HO1	LEU	63	8.237	-
13.610	1.532	1.00	0.00	BD ATOM	862	HO2	ARG	51	2.206	4.515	-12.715	1.00	0.00	BD ATOM	956	HO2	LEU	57	-6.778	2.962	-6.442	1.00	0.00	BD ATOM	1050	HO1	LEU	63	8.237	-
13.610	1.532	1.00	0.00	BD ATOM	863	HO2	ARG	51	2.206	4.515	-12.715	1.00	0.00	BD ATOM	957	HO2	LEU													

REPLACEMENT SHEET

6.329	-9.898	1.00	0.00	BrD ATOM	1099	NE	ANG	66	13.162	-	2.400	BrD ATOM	1287	C	ALA	76	2.240
5.351	-10.237	1.00	0.00	BrD ATOM	1101	CE	ANG	66	13.688	-	4.084	BrD ATOM	1199	CE	ALA	76	1.447
5.255	-11.059	1.00	0.00	BrD ATOM	1102	HN	ANG	66	13.801	-	4.316	BrD ATOM	1195	HN	ALA	71	2.121
5.407	-9.074	1.00	0.00	BrD ATOM	1103	HN1	ANG	66	13.117	-	3.879	BrD ATOM	1197	HN2	LVS	71	13.887
5.726	-7.965	1.00	0.00	BrD ATOM	1104	HN2	ANG	66	15.599	-	3.334	BrD ATOM	1198	HN3	LVS	71	14.334
5.726	-7.965	1.00	0.00	BrD ATOM	1105	HN3	ANG	66	15.599	-	4.880	BrD ATOM	1199	HN4	LVS	71	15.252
5.593	-7.965	1.00	0.00	BrD ATOM	1106	HN4	ANG	66	15.599	-	3.100	BrD ATOM	1200	HN5	LVS	71	9.041
5.334	-9.037	1.00	0.00	BrD ATOM	1107	HN5	ANG	66	17.604	-	2.529	BrD ATOM	1201	HN6	LVS	72	9.076
5.327	-8.161	1.00	0.00	BrD ATOM	1108	HN6	ANG	66	17.604	-	4.447	BrD ATOM	1202	HN7	LVS	72	9.933
5.237	-9.887	1.00	0.00	BrD ATOM	1109	HN7	ANG	66	10.708	-	4.888	BrD ATOM	1203	HN8	LVS	72	7.844
1.474	-10.682	1.00	0.00	BrD ATOM	1110	HN8	ANG	66	11.954	-	5.201	BrD ATOM	1204	HN9	LVS	72	7.366
0.405	-10.956	1.00	0.00	BrD ATOM	1111	HN9	ANG	66	8.346	-	4.818	BrD ATOM	1205	HN10	LVS	72	8.167
1.159	-10.631	1.00	0.00	BrD ATOM	1112	HN10	ANG	66	9.716	-	7.212	BrD ATOM	1206	HN11	LVS	72	7.249
0.780	-11.472	1.00	0.00	BrD ATOM	1113	HN11	ANG	66	9.952	-	7.070	BrD ATOM	1207	HN12	LVS	72	9.531
0.105	-12.164	1.00	0.00	BrD ATOM	1114	HN12	ANG	66	7.982	-	0.364	BrD ATOM	1208	HN13	LVS	72	9.870
0.065	-11.738	1.00	0.00	BrD ATOM	1115	HN13	ANG	66	7.062	-	6.150	BrD ATOM	1209	HN14	LVS	72	9.629
0.230	-12.984	1.00	0.00	BrD ATOM	1116	HN14	ANG	66	8.276	-	6.588	BrD ATOM	1210	HN15	LVS	72	8.419
2.639	-11.976	1.00	0.00	BrD ATOM	1117	HN15	ANG	66	7.713	-	7.004	BrD ATOM	1211	HN16	LVS	72	8.419
2.385	-10.973	1.00	0.00	BrD ATOM	1118	HN16	ANG	66	7.042	-	7.164	BrD ATOM	1212	HN17	LVS	72	9.135
2.032	-14.005	1.00	0.00	BrD ATOM	1119	HN17	ANG	66	8.125	-	7.164	BrD ATOM	1213	HN18	LVS	72	9.135
1.301	-14.604	1.00	0.00	BrD ATOM	1120	HN18	ANG	66	8.645	-	8.208	BrD ATOM	1214	HN19	LVS	72	7.758
1.899	-12.475	1.00	0.00	BrD ATOM	1121	HN19	ANG	66	6.787	-	8.869	BrD ATOM	1215	HN20	LVS	72	6.474
3.425	-11.872	1.00	0.00	BrD ATOM	1122	HN20	ANG	66	7.872	-	7.868	BrD ATOM	1216	HN21	LVS	72	8.150
3.403	-11.872	1.00	0.00	BrD ATOM	1123	HN21	ANG	66	7.872	-	8.956	BrD ATOM	1217	HN22	LVS	72	8.150
3.541	-15.516	1.00	0.00	BrD ATOM	1124	HN22	ANG	66	9.198	-	8.956	BrD ATOM	1218	HN23	LVS	72	6.758
4.320	-13.749	1.00	0.00	BrD ATOM	1125	HN23	ANG	66	6.531	-	9.266	BrD ATOM	1219	HN24	LVS	72	6.882
5.475	-14.255	1.00	0.00	BrD ATOM	1126	HN24	ANG	66	6.531	-	5.034	BrD ATOM	1220	HN25	LVS	72	7.370
6.127	-13.753	1.00	0.00	BrD ATOM	1127	HN25	ANG	66	7.446	-	5.001	BrD ATOM	1221	HN26	LVS	72	8.343
1.006	-9.840	1.00	0.00	BrD ATOM	1128	HN26	ANG	66	8.910	-	4.933	BrD ATOM	1222	HN27	LVS	72	6.528
0.181	-9.839	1.00	0.00	BrD ATOM	1129	HN27	ANG	66	7.880	-	4.786	BrD ATOM	1223	HN28	LVS	72	7.982
0.647	-9.193	1.00	0.00	BrD ATOM	1130	HN28	ANG	66	7.686	-	3.728	BrD ATOM	1224	HN29	LVS	72	7.394
0.855	-7.736	1.00	0.00	BrD ATOM	1131	HN29	ANG	66	7.528	-	3.478	BrD ATOM	1225	HN30	LVS	72	7.447
1.924	-7.788	1.00	0.00	BrD ATOM	1132	HN30	ANG	66	5.918	-	4.596	BrD ATOM	1226	HN31	LVS	72	6.886
0.368	-8.188	1.00	0.00	BrD ATOM	1133	HN31	ANG	66	5.664	-	4.205	BrD ATOM	1227	HN32	LVS	72	5.937
2.734	-6.512	1.00	0.00	BrD ATOM	1134	HN32	ANG	66	4.987	-	4.787	BrD ATOM	1228	HN33	LVS	72	8.166
2.712	-6.664	1.00	0.00	BrD ATOM	1135	HN33	ANG	66	5.378	-	3.997	BrD ATOM	1229	HN34	LVS	72	8.729
1.732	-9.101	1.00	0.00	BrD ATOM	1136	HN34	ANG	66	3.764	-	3.507	BrD ATOM	1230	HN35	LVS	72	7.512
0.927	-6.638	1.00	0.00	BrD ATOM	1137	HN35	ANG	66	3.764	-	5.701	BrD ATOM	1231	HN36	LVS	72	7.104
3.788	-8.933	1.00	0.00	BrD ATOM	1138	HN36	ANG	66	4.526	-	6.707	BrD ATOM	1232	HN37	LVS	72	5.659
4.591	-9.979	1.00	0.00	BrD ATOM	1139	HN37	ANG	66	2.855	-	6.931	BrD ATOM	1233	HN38	LVS	72	7.237
2.797	-7.110	1.00	0.00	BrD ATOM	1140	HN38	ANG	66	1.868	-	7.243	BrD ATOM	1234	HN39	LVS	72	5.528
3.807	-8.374	1.00	0.00	BrD ATOM	1141	HN39	ANG	66	2.416	-	3.665	BrD ATOM	1235	HN40	LVS	72	6.413
4.855	-8.639	1.00	0.00	BrD ATOM	1142	HN40	ANG	66	8.206	-	2.432	BrD ATOM	1236	HN41	LVS	72	6.351
5.490	-7.920	1.00	0.00	BrD ATOM	1143	HN41	ANG	66	8.001	-	1.253	BrD ATOM	1237	HN42	LVS	72	4.364
0.327	-6.548	1.00	0.00	BrD ATOM	1144	HN42	ANG	66	9.149	-	1.359	BrD ATOM	1238	HN43	LVS	72	5.984
0.750	-5.416	1.00	0.00	BrD ATOM	1145	HN43	ANG	66	9.268	-	0.003	BrD ATOM	1239	HN44	LVS	72	6.740
0.888	-7.735	1.00	0.00	BrD ATOM	1146	HN44	ANG	66	9.137	-	0.181	BrD ATOM	1240	HN45	LVS	72	6.413
1.170	-5.789	1.00	0.00	BrD ATOM	1147	HN45	ANG	66	11.231	-	0.181	BrD ATOM	1241	HN46	LVS	72	6.413
1.941	-5.789	1.00	0.00	BrD ATOM	1148	HN46	ANG	66	11.231	-	0.181	BrD ATOM	1242	HN47	LVS	72	6.413
2.512	-5.291	1.00	0.00	BrD ATOM	1149	HN47	ANG	66	11.389	-	1.240	BrD ATOM	1243	HN48	LVS	72	4.364
1.146	-7.146	1.00	0.00	BrD ATOM	1150	HN48	ANG	66	12.954	-	1.851	BrD ATOM	1244	HN49	LVS	72	5.412
2.937	-7.117	1.00	0.00	BrD ATOM	1151	HN49	ANG	66	12.107	-	1.451	BrD ATOM	1245	HN50	LVS	72	4.546
0.718	-7.032	1.00	0.00	BrD ATOM	1152	HN50	ANG	66	11.701	-	1.797	BrD ATOM	1246	HN51	LVS	72	4.691
0.718	-7.032	1.00	0.00	BrD ATOM	1153	HN51	ANG	66	11.701	-	1.332	BrD ATOM	1247	HN52	LVS	72	4.181
0.166	-6.420	1.00	0.00	BrD ATOM	1154	HN52	ANG	66	12.021	-	2.953	BrD ATOM	1248	HN53	LVS	72	4.039
0.111	-6.420	1.00	0.00	BrD ATOM	1155	HN53	ANG	66	11.389	-	2.953	BrD ATOM	1249	HN54	LVS	72	3.780
2.572	-4.899	1.00	0.00	BrD ATOM	1156	HN54	ANG	66	12.954	-	2.942	BrD ATOM	1250	HN55	LVS	72	3.593
1.889	-4.494	1.00	0.00	BrD ATOM	1157	HN55	ANG	66	12.954	-	3.362	BrD ATOM	1251	HN56	LVS	72	3.593
2.983	-5.716	1.00	0.00	BrD ATOM	1158	HN56	ANG	66	10.409	-	3.538	BrD ATOM	1252	HN57	LVS	72	2.989
0.506	-1.565	1.00	0.00	BrD ATOM	1159	HN57	ANG	66	10.409	-	1.116	BrD ATOM	1253	HN58	LVS	72	4.643
1.132	-5.110	1.00	0.00	BrD ATOM	1160	HN58	ANG	66	10.501	-	0.694	BrD ATOM	1254	HN59	LVS	72	3.503
1.306	-6.044	1.00	0.00	BrD ATOM	1161	HN59	ANG	66	10.501	-	0.694	BrD ATOM	1255	HN60	LVS	72	3.503
5.439	-4.088	1.00	0.00	BrD ATOM	1162	HN60	ANG	66	10.268	-	1.755	BrD ATOM	1256	HN61	LVS	72	5.612
2.190	-4.189	1.00	0.00	BrD ATOM	1163	HN61	ANG	66	10.593	-	0.403	BrD ATOM	1257	HN62	LVS	72	4.605
2.146	-4.081	1.00	0.00	BrD ATOM	1164	HN62	ANG	66	11.246	-	1.308	BrD ATOM	1258	HN63	LVS	72	6.100
3.552	-4.767	1.00	0.00	BrD ATOM	1165	HN63	ANG	66	9.546	-	0.744	BrD ATOM	1259	HN64	LVS	72	7.021
3.183	-5.218	1.00	0.00	BrD ATOM	1166	HN64	ANG	66	10.488	-	0.029	BrD ATOM	1260	HN65	LVS	72	7.324
4.547	-3.758	1.00	0.00	BrD ATOM	1167	HN65	ANG	66	10.268	-	0.029	BrD ATOM	1261	HN66	LVS	72	8.086
5.439	-4.088	1.00	0.00	BrD ATOM	1168	HN66	ANG	66	10.268	-	0.744	BrD ATOM	1262	HN67	LVS	72	8.173
2.011	-2.810	1.00	0.00	BrD ATOM	1169	HN67	ANG	66	10.881	-	0.403	BrD ATOM	1263	HN68	LVS	72	7.211
1.306	-2.674	1.00	0.00	BrD ATOM	1170	HN68	ANG	66	11.039	-	0.049	BrD ATOM	1264	HN69	LVS	72	8.173
2.605	-1.791	1.00	0.00	BrD ATOM	1171	HN69	ANG	66	11.039	-	2.957	BrD ATOM	1265	HN70	LVS	72	8.173
2.470	-0.432	1.00	0.00	BrD ATOM	1172	HN70	ANG	66	11.419	-	2.415	BrD ATOM	1266	HN71	LVS	72	8.173
1.415	-0.226	1.00	0.00	BrD ATOM	1173	HN71	ANG	66	11.419	-	2.415	BrD ATOM	1267	HN72	LVS	72	8.173
3.036	-0.564	1.00	0.00	BrD ATOM	1174	HN72	ANG	66	10.915	-	2.914	BrD ATOM	1268	HN73	LVS	72	4.312
3.931	-0.162	1.00	0.00	BrD ATOM	1175	HN73	ANG	66	11.848	-	2.052	BrD ATOM	1269	HN74	LVS	72	4.312
3.931	-0.162	1.00	0.00	BrD ATOM	1176	HN74	ANG	66	12.550	-	3.642	BrD ATOM	1270	HN75	LVS	72	4.312
2.261	-0.191	1.00	0.00	BrD ATOM	1177	HN75	ANG	66	13.139	-	4.814	BrD ATOM	1271	HN76	LVS	72	4.312
2.261	-0.191	1.00	0.00	BrD ATOM	1178	HN76	ANG	66	13.139	-	4.814	BrD ATOM	12				

0.514	3.764	1.00	0.00	0.00	BD ATOM	1381	CB	PHE	82	-4.131	-	5.541	4.446	1.00	0.00	0.00	BD ATOM	1475	CA	TYR	88	-12.759	-	7.446	4.078	1.00	0.00	0.00	BD ATOM	1569	OEI	GLU	94	-16.617	-
0.333	4.699	1.00	0.00	0.00	BD ATOM	1382	HEB	PHE	82	-4.466	-	6.319	4.446	1.00	0.00	0.00	BD ATOM	1476	HA	TYR	88	-13.070	-	7.446	4.078	1.00	0.00	0.00	BD ATOM	1570	OEI	GLU	94	-16.617	-
0.220	5.511	1.00	0.00	0.00	BD ATOM	1383	HEB	PHE	82	-4.712	-	7.330	4.344	1.00	0.00	0.00	BD ATOM	1477	CB	TYR	88	-12.361	-	8.370	3.840	1.00	0.00	0.00	BD ATOM	1571	O	GLU	94	-14.321	-
1.484	5.536	1.00	0.00	0.00	BD ATOM	1384	CO	PHE	82	-3.261	-	6.330	2.568	1.00	0.00	0.00	BD ATOM	1478	HEB	TYR	88	-11.310	-	3.556	5.725	1.00	0.00	0.00	BD ATOM	1572	O	GLU	94	-14.321	-
1.959	4.028	1.00	0.00	0.00	BD ATOM	1385	COI	PHE	82	-1.979	-	6.502	2.489	1.00	0.00	0.00	BD ATOM	1479	HEB	TYR	88	-12.537	-	4.137	5.876	1.00	0.00	0.00	BD ATOM	1573	N	TYR	95	-15.387	-
1.977	5.372	1.00	0.00	0.00	BD ATOM	1386	HEB	PHE	82	-1.606	-	5.588	2.139	1.00	0.00	0.00	BD ATOM	1480	COI	TYR	88	-13.128	-	2.357	5.150	1.00	0.00	0.00	BD ATOM	1574	HA	TYR	95	-15.387	-
2.162	2.809	1.00	0.00	0.00	BD ATOM	1387	HEB	PHE	82	-4.723	-	8.266	1.905	1.00	0.00	0.00	BD ATOM	1481	COI	TYR	88	-12.606	-	1.930	5.057	1.00	0.00	0.00	BD ATOM	1575	CA	TYR	95	-13.927	-
2.827	2.487	1.00	0.00	0.00	BD ATOM	1388	HEB	PHE	82	-1.177	-	8.896	1.914	1.00	0.00	0.00	BD ATOM	1482	COI	TYR	88	-11.372	-	2.327	3.840	1.00	0.00	0.00	BD ATOM	1576	HA	TYR	95	-14.448	-
3.966	6.982	1.00	0.00	0.00	BD ATOM	1389	HEB	PHE	82	-0.179	-	7.066	1.213	1.00	0.00	0.00	BD ATOM	1483	HEB	TYR	88	-13.792	-	0.260	4.165	1.00	0.00	0.00	BD ATOM	1577	HEB	TYR	95	-14.448	-
2.952	3.951	1.00	0.00	0.00	BD ATOM	1390	CE2	PHE	82	-2.928	-	6.087	1.392	1.00	0.00	0.00	BD ATOM	1484	HEB	TYR	88	-12.880	-	0.160	4.827	1.00	0.00	0.00	BD ATOM	1578	HEB	TYR	95	-14.448	-
2.952	2.004	1.00	0.00	0.00	BD ATOM	1391	CE2	PHE	82	-3.302	-	9.568	0.755	1.00	0.00	0.00	BD ATOM	1485	CE2	TYR	88	-15.075	-	0.367	4.176	1.00	0.00	0.00	BD ATOM	1579	HEB	TYR	95	-14.270	-
3.284	1.073	1.00	0.00	0.00	BD ATOM	1392	CE2	PHE	82	-1.651	-	10.562	0.578	1.00	0.00	0.00	BD ATOM	1486	CE2	TYR	88	-16.042	-	1.350	2.245	1.00	0.00	0.00	BD ATOM	1580	CE2	TYR	95	-15.041	-
3.284	1.073	1.00	0.00	0.00	BD ATOM	1393	CE2	PHE	82	-1.651	-	7.959	0.462	1.00	0.00	0.00	BD ATOM	1487	CE2	TYR	88	-16.042	-	0.367	4.176	1.00	0.00	0.00	BD ATOM	1581	CE2	TYR	95	-15.041	-
3.906	1.709	1.00	0.00	0.00	BD ATOM	1394	CE2	PHE	82	-6.745	-	9.238	0.216	1.00	0.00	0.00	BD ATOM	1488	CE2	TYR	88	-16.042	-	0.367	4.176	1.00	0.00	0.00	BD ATOM	1582	CE2	TYR	95	-15.041	-
1.197	4.994	1.00	0.00	0.00	BD ATOM	1395	O	PHE	82	-6.745	-	9.238	0.216	1.00	0.00	0.00	BD ATOM	1489	CE2	TYR	88	-16.042	-	0.367	4.176	1.00	0.00	0.00	BD ATOM	1583	CE2	TYR	95	-15.041	-
2.185	5.431	1.00	0.00	0.00	BD ATOM	1396	O	PHE	82	-6.745	-	10.145	-0.512	1.00	0.00	0.00	BD ATOM	1490	CE2	TYR	88	-13.993	-	1.377	0.963	1.00	0.00	0.00	BD ATOM	1584	CE2	TYR	95	-15.951	-
2.185	5.241	1.00	0.00	0.00	BD ATOM	1397	N	TYR	83	-3.841	-	10.465	-0.014	1.00	0.00	0.00	BD ATOM	1491	HE	TYR	88	-14.309	-	2.208	0.579	1.00	0.00	0.00	BD ATOM	1585	HE	TYR	95	-14.656	-
2.149	4.852	1.00	0.00	0.00	BD ATOM	1398	HA	TYR	83	-5.427	-	4.711	4.256	1.00	0.00	0.00	BD ATOM	1492	O	TYR	88	-14.309	-	0.901	0.673	1.00	0.00	0.00	BD ATOM	1586	HE2	TYR	95	-14.546	-
2.337	6.048	1.00	0.00	0.00	BD ATOM	1399	HA	TYR	83	-5.427	-	4.711	4.256	1.00	0.00	0.00	BD ATOM	1493	N	ASN	89	-14.012	-	1.785	0.062	1.00	0.00	0.00	BD ATOM	1587	HE2	TYR	95	-15.131	-
2.337	6.048	1.00	0.00	0.00	BD ATOM	1400	HA	TYR	83	-5.427	-	4.711	4.256	1.00	0.00	0.00	BD ATOM	1494	N	ASN	89	-14.012	-	0.210	0.182	1.00	0.00	0.00	BD ATOM	1588	HE2	TYR	95	-15.131	-
4.512	6.216	1.00	0.00	0.00	BD ATOM	1401	HE	TYR	83	-3.927	-	5.288	4.482	1.00	0.00	0.00	BD ATOM	1495	HA	ASN	89	-13.518	-	0.508	1.079	1.00	0.00	0.00	BD ATOM	1589	HE2	TYR	95	-15.131	-
4.512	6.216	1.00	0.00	0.00	BD ATOM	1402	HE	TYR	83	-3.927	-	5.288	4.482	1.00	0.00	0.00	BD ATOM	1496	HA	ASN	89	-13.518	-	0.508	1.079	1.00	0.00	0.00	BD ATOM	1590	HE2	TYR	95	-15.131	-
3.666	6.982	1.00	0.00	0.00	BD ATOM	1403	COI	TYR	83	-4.138	-	4.436	4.956	1.00	0.00	0.00	BD ATOM	1497	HEB	ASN	89	-15.816	-	1.552	5.763	1.00	0.00	0.00	BD ATOM	1591	HEB	ASN	96	-12.733	-
5.589	7.913	1.00	0.00	0.00	BD ATOM	1404	COI	TYR	83	-4.283	-	3.023	6.192	1.00	0.00	0.00	BD ATOM	1498	HEB	ASN	89	-15.816	-	1.552	5.763	1.00	0.00	0.00	BD ATOM	1592	HEB	ASN	96	-12.733	-
6.222	6.192	1.00	0.00	0.00	BD ATOM	1405	COI	TYR	83	-4.283	-	2.421	6.597	1.00	0.00	0.00	BD ATOM	1499	HEB	ASN	89	-15.816	-	1.552	5.763	1.00	0.00	0.00	BD ATOM	1593	HEB	ASN	96	-12.733	-
6.222	6.192	1.00	0.00	0.00	BD ATOM	1406	COI	TYR	83	-4.283	-	2.421	6.597	1.00	0.00	0.00	BD ATOM	1500	HEB	ASN	89	-15.816	-	1.552	5.763	1.00	0.00	0.00	BD ATOM	1594	HEB	ASN	96	-12.733	-
6.222	6.192	1.00	0.00	0.00	BD ATOM	1407	COI	TYR	83	-4.283	-	2.421	6.597	1.00	0.00	0.00	BD ATOM	1501	HEB	ASN	89	-15.816	-	1.552	5.763	1.00	0.00	0.00	BD ATOM	1595	HEB	ASN	96	-12.733	-
6.222	6.192	1.00	0.00	0.00	BD ATOM	1408	COI	TYR	83	-4.283	-	2.421	6.597	1.00	0.00	0.00	BD ATOM	1502	HEB	ASN	89	-15.816	-	1.552	5.763	1.00	0.00	0.00	BD ATOM	1596	HEB	ASN	96	-12.733	-
5.391	7.749	1.00	0.00	0.00	BD ATOM	1409	COI	TYR	83	-4.283	-	2.421	6.597	1.00	0.00	0.00	BD ATOM	1503	HEB	ASN	89	-15.816	-	1.552	5.763	1.00	0.00	0.00	BD ATOM	1597	HEB	ASN	96	-12.733	-
4.077	5.402	1.00	0.00	0.00	BD ATOM	1410	O	TYR	83	-7.073	-	1.906	7.755	1.00	0.00	0.00	BD ATOM	1504	COI	ASN	89	-13.444	-	0.860	9.365	1.00	0.00	0.00	BD ATOM	1598	HEB	TYR	96	-11.431	-
4.735	6.082	1.00	0.00	0.00	BD ATOM	1411	N	ASN	84	-6.391	-	1.099	7.336	1.00	0.00	0.00	BD ATOM	1505	COI	ASN	89	-12.731	-	1.522	9.615	1.00	0.00	0.00	BD ATOM	1599	HEB	TYR	96	-11.431	-
3.956	4.082	1.00	0.00	0.00	BD ATOM	1412	HA	ASN	84	-5.732	-	1.909	6.347	1.00	0.00	0.00	BD ATOM	1506	C	ASN	89	-16.332	-	0.114	9.116	1.00	0.00	0.00	BD ATOM	1600	HE2	TYR	96	-12.977	-
3.418	3.597	1.00	0.00	0.00	BD ATOM	1413	CA	ASN	84	-7.458	-	4.982	6.985	1.00	0.00	0.00	BD ATOM	1507	O	ASN	89	-16.088	-	0.708	10.594	1.00	0.00	0.00	BD ATOM	1601	COI	TYR	96	-10.351	-
4.609	3.354	1.00	0.00	0.00	BD ATOM	1414	HA	ASN	84	-7.458	-	4.982	6.985	1.00	0.00	0.00	BD ATOM	1508	N	ASN	89	-17.296	-	0.458	10.477	1.00	0.00	0.00	BD ATOM	1602	COI	TYR	96	-10.351	-
4.609	3.354	1.00	0.00	0.00	BD ATOM	1415	HA	ASN	84	-7.458	-	4.982	6.985	1.00	0.00	0.00	BD ATOM	1509	N	ASN	89	-17.296	-	0.458	10.477	1.00	0.00	0.00	BD ATOM	1603	COI	TYR	96	-10.351	-
4.609	3.354	1.00	0.00	0.00	BD ATOM	1416	HA	ASN	84	-7.458	-	4.982	6.985	1.00	0.00	0.00	BD ATOM	1510	N	ASN	89	-17.296	-	0.458	10.477	1.00	0.00	0.00	BD ATOM	1604	COI	TYR	96	-10.351	-
3.629	1.464	1.00	0.00	0.00	BD ATOM	1417	HEB	ASN	84	-6.118	-	6.058	7.470	1.00	0.00	0.00	BD ATOM	1511	COI	ASN	89	-18.442	-	0.375	9.493	1.00	0.00	0.00	BD ATOM	1605	COI	TYR	96	-12.252	-
5.086	1.716	1.00	0.00	0.00	BD ATOM	1418	COI	ASN	84	-8.115	-	7.122	8.258	1.00	0.00	0.00	BD ATOM	1512	COI	ASN	89	-19.228	-	1.008	11.979	1.00	0.00	0.00	BD ATOM	1606	COI	TYR	96	-12.252	-
5.086	1.716	1.00	0.00	0.00	BD ATOM	1419	COI	ASN	84	-8.115	-	7.291	7.006	1.00	0.00	0.00	BD ATOM	1513	HEB	ALA	90	-19.280	-	0.319	11.594	1.00	0.00	0.00	BD ATOM	1607	HEB	TYR	96	-8.495	-
5.946	-0.113	1.00	0.00	0.00	BD ATOM	1420	NO2	ASN	84	-7.677	-	7.045	5.956	1.00	0.00	0.00	BD ATOM	1514	HEB	ALA	90	-18.997	-	0.132	11.480	1.00	0.00	0.00	BD ATOM	1608	HEB	TYR	96	-11.458	-
5.946	-0.113	1.00	0.00	0.00	BD ATOM	1421	NO2	ASN	84	-7.677	-	7.045	5.956	1.00	0.00																				

REPLACEMENT SHEET

4.534	5.217	1.00	0.00	BRD	1945	CDL	11E	116	4.662
0.615	6.373	1.00	0.00	BRD	1946	HD11	11E	116	3.870
0.073	5.875	1.00	0.00	BRD	1947	HD12	11E	116	4.240
1.328	7.147	1.00	0.00	BRD	1948	HD13	11E	116	5.356
1.481	8.419	1.00	0.00	BRD	1949	HD14	11E	116	5.802
1.751	8.053	1.00	0.00	BRD	1950	O	11E	116	5.802
0.544	8.154	1.00	0.00	BRD	1951	N	ASP	117	6.222
2.465	9.014	1.00	0.00	BRD	1952	HN	ASP	117	6.079
3.424	8.874	1.00	0.00	BRD	1953	CA	ASP	117	5.805
1.862	10.275	1.00	0.00	BRD	1954	HA	ASP	117	5.197
2.985	10.798	1.00	0.00	BRD	1955	CB	ASP	117	4.967
3.424	10.798	1.00	0.00	BRD	1956	HD1	ASP	117	5.564
0.267	10.220	1.00	0.00	BRD	1957	HD2	ASP	117	4.168
0.596	8.976	1.00	0.00	BRD	1958	CG	ASP	117	3.722
0.555	10.876	1.00	0.00	BRD	1959	OD1	ASP	117	2.897
1.488	10.787	1.00	0.00	BRD	1960	OD2	ASP	117	3.572
0.421	11.641	1.00	0.00	BRD	1961	C	ASP	117	7.006
1.305	11.154	1.00	0.00	BRD	1962	O	ASP	117	6.846
1.328	11.154	1.00	0.00	BRD	1963	N	ASP	118	8.205
1.897	10.725	1.00	0.00	BRD	1964	HN	LVS	118	9.187
2.384	9.879	1.00	0.00	BRD	1965	CA	LVS	118	9.416
1.613	11.488	1.00	0.00	BRD	1966	HA	LVS	118	10.248
2.063	10.968	1.00	0.00	BRD	1967	CB	LVS	118	9.315
2.220	12.889	1.00	0.00	BRD	1968	HB1	LVS	118	8.636
1.624	13.480	1.00	0.00	BRD	1969	HD2	LVS	118	10.292
3.424	13.480	1.00	0.00	BRD	1970	CG	LVS	118	8.018
3.656	12.895	1.00	0.00	BRD	1971	HD3	LVS	118	7.271
3.719	12.296	1.00	0.00	BRD	1972	HD2	LVS	118	8.595
3.946	13.911	1.00	0.00	BRD	1973	CD	LVS	118	9.857
4.608	12.327	1.00	0.00	BRD	1974	HD1	LVS	118	10.477
4.972	13.133	1.00	0.00	BRD	1975	HD2	LVS	118	10.468
4.073	11.613	1.00	0.00	BRD	1976	CE	LVS	118	9.208
5.725	11.903	1.00	0.00	BRD	1977	HB1	LVS	118	8.502
5.725	12.903	1.00	0.00	BRD	1978	HB2	LVS	118	8.502
6.386	12.369	1.00	0.00	BRD	1979	NZ	LVS	118	10.212
6.652	10.943	1.00	0.00	BRD	1980	HD1	LVS	118	11.163
6.458	11.316	1.00	0.00	BRD	1981	HD2	LVS	118	9.989
7.656	11.096	1.00	0.00	BRD	1982	HD3	LVS	118	10.206
6.459	9.921	1.00	0.00	BRD	1983	C	LVS	118	9.657
0.410	11.501	1.00	0.00	BRD	1984	OT1	LVS	118	9.047
0.420	11.501	1.00	0.00	BRD	1985	OT2	LVS	118	10.454
0.302	12.459	1.00	0.00	BRD	END				-